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ABQ75387/c
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                                                              The invention describes a compound (I) 8-50 nucleobases in length targeted to a nucleic acid molecule (II) encoding RAIDD which is an adaptor molecule containing both death domain (DD) and caspase recruitment domains (CARD), where (I) specifically hybridises with and inhibits expression of RAIDD, or specifically hybridises with at least an encleobase portion of an active site on (II). (I) is useful for inhibiting the expression of RAIDD (Receptor interacting protein (RIP) associated ICH-I/CED-3-homologous protein with death domain) in cells or inhibiting the for treating an animal having a disease or condition associated with RAIDD, where the disease or condition is a hyperproliferative disorder such as cancer, or a growth or metabolic disorder. (I) is also useful for distinguishing functions of various members of a biological pathway, and in antisense gene therapy. (I) is also susful prophylactically, e.g. to prevent or delay infection, inflammation or tumour formation. This sequence represents a mouse RAIDD antisense oligonucleotide used to control expression of the RAIDD protein
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encoding RAIDD which is an adaptor molecule containing both death domain and caspase recruitment domains, for treating hyperproliferative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /note= "2'-0-methoxyethyl gapmer with an 8 nucleotide deoxy gap and a phosphorothioate backbone; cytosine residues are 5-methyl cytosines"
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                                                                                                                                                                                                                                                                                                0.8%; Score 14.2; DB 1; Length 20; 34.2%; Pred. No. 8.7e+02;
                                                                                                                                                                                                                                                                                                                      3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human RNase HII antisense oligonucleotide SEQ ID NO:20.
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                                                                                                                                                                                                                                                                                                                      Mismatches
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                                             Claim 3; Page 94; 144pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             base= OTHER
                                                                                                                                                                                                                                                                                                                                             54
                                                                                                                                                                                                                                                                                                                                                                 1 GAAGGCAGGATGTCCAGCA 19
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es 16; Conservative
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Matches
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with a disease or condition associated with a human RNase H, for inhibiting the expression of a protein, or for reducing cellular RNA via antisense technology.
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14.2%; Pred. No. 8.7e+02;
ve 0; Mismatches 3; Indels
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                                                                                                                                                   Claim 38; Page 37; 70pp; English.
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RESULT 1029
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                                  The present invention describes a method for promoting the inhibition of the expression of a protein comprising employing a mammalian RNase H polypeptide so that cleavage of an RN Strand of an oligonucleotide-RNA complex duplex occurs. Also described is a compound 8 to 50 nucleobases in length targeted to the nucleic acid encoding the human RNase HII polypeptide, where the compound specifically hybridises with and inhibits the expression of a human RNase HII polypeptide is useful for inhibiting the expression of an antisense oligonucleotide, is useful for inhibiting the expression of treating an animal with a disease or condition associated with a human RNase HII polypeptide in the sacciated with a human RNase HII polypeptide. The method is useful for inhibiting the expression of a protein, particularly for reducing callular RNA via antisense technology. The present sequence represents a human RNase HII antisense oligonucleotide, which is used in an example from the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The present sequence represents an oligonucleotide which targets polymucleotides encoding human aurora 2 kinase. The oligonucleotide inhibits aurora 2 kinase expression. The oligonucleotide is useful in the diagnosis and treatment of cancers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
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Claim 38; Page 37; 70pp; English.
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The invention relates to a map of the bread wheat D genome comprising the genome location of a microsatellite markers selected from a group of 185 such markers (A8022913-A802917). The invention also encompasses the use of left (A8022918-A8023107) and right (A8023103-A802327) primers to of left (A8022918-A8023107) and right (A8023103-A802327) primers to complete the microsatellite markers and to identify genes responsible for a phenotypic trait of interest in wheat. Wheat is an allohexaploid species consisting of 3 diploid genomes designated A, B and C responsible for a phenotypic trait of interest in wheat. Wheat is an office allohexaploid species consisting of 3 diploid genomes designated A, B and C b, resulting from two successive intercrossings involving at least three different species. The D genome is thought to have been introduced in the most recent intercrossing, between the ampliploid AABB and Triticum tauschil (DD), probably involving only a limited number of genome, and its low level of polymorphism, the genetic mapping of wheat has to date been difficult. Microsatellites are tandemely repeated sequences between one and six nuclectides long, and are very polymorphic in length contracted to polymorphism makes them especially suitable for the genetic mapping of species which show little intraspecies polymorphism, such as wheat. In addition, microsatellite anterest of the invention are comminged from the ancestral diploid donor species Triticum tauschil and genomes. These microsatellite markers thus help to overcome some of the problems associated with the genetic mapping of wheat. The wheat D genome of the invention are useful for identifying genes responsible for a phenotypic canneaution are useful for identifying genes responsible for a phenotypic particular they may be used for analysing genes and alleles implicated in disease and for identifying development factors, quality factors and conferring resistance to pathogens and xenobictics. The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Map of wheat D genome comprising the genome location of a microsatellite marker, useful for e.g. identifying genes responsible for a desired phenotypic trait, especially quantitative trait loci in wheat, and
                                                                                                                                                             T. tauschii/wheat D genome microsatellite cfd226 right PCR primer.
                                                                                                                                                                                                      Microsatellite marker, wheat, D genome, mapping; genotyping, polymorphism, phenotypic trait, QTL; quantitative trait locus; disease-associated gene; development factor; quality factor; resistance factor; wheat product, identification; detection; genetically modified wheat; PCR; primer; 88.
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ABQ93219 standard; DNA; 20 BP.
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                                                                                            (revised)
(first entry)
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                                                                                                                                                                                                                                                                                                                                                 Aegilops tauschii.
Triticum aestivum.
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                                                                                          29-AUG-2003
21-OCT-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The present invention describes an isolated peptide (I) consisting of an amino acid sequence selected from: (a) the amino acid sequence of a variant of the oestrogen receptor alpha (ESR-alpha) protein in AAG68251; or (b) a fragment comprising at least 10 contiguous amino acids of the protein in AAG68251. (I) has cytostatic, osteopathic, cardiant and vasctropic activities, and can be used in gene therapy and vaccine production. (I) is useful for identifying an agent that binds to (I), by contacting (I) with an agent and assaying the contacted mixture to determine whether a complex is formed with the agent bound to the peptide. A polynucleotide (II), encoding (I), is useful in the development of diagnostics and therapies for diseases and disorders
          mapping and genotyping diploid and polyploid species of Triticum, particularly Aegilops, Triticum monococcum, Triticum durum, Triticum aestivum, or related species; for identifying cultivars and hybrids of Triticum and related species; for assess whether or not a product comprises wheat or a related species; and to assess whether or not a product comprises wheat or a related species; and to assess whether or not a product comprises genetically modified wheat. The present sequence represents a specifically claimed Triticum tauschii/wheat genome D microsatellite marker right PCR primer of the invention. (Updated on 29-MUG-2003 to standardise OS fleid)
                                                                                                                                                                                                                                                                                                                                                                                                                                         Human, oestrogen receptor alpha, ESR-alpha, ER, chromosome 6; Syne-2; synaptic nuclei expressed gene 2; haplotype; cytostatic; osteopathic; cardiant; vasotropic; gene therapy; vaccine; cancer; osteoporosis; cardiant; vasotropic, eservogen receptor; PCR primer; sequencing; ss. cardiavascular disease; oestrogen receptor; PCR primer; sequencing; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Novel variant of estrogen receptor alpha polypeptide useful for determining the biological activity of a protein for high throughput screening and for raising antibodies that elicit an immune response in
and associated primers may be also be used in
                                                                                                                                                                                                           Gaps
                                                                                                                                                                                                           ó,
                                                                                                                                                                            Score 14.2; DB 1; Length 20;
Pred. No. 8.7e+02;
0; Mismatches 3; Indels
                                                                                                                                                 Sequence 20 BP; 4 A; 5 C; 4 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Winn-Deen ES;
                                                                                                                                                                                                                                                                                                                                                                                                                 Destrogen receptor alpha gene PCR primer #14.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Hwang SS,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 17; Fig 2c; 333pp; English.
                                                                                                                                                                                                                                    792 CGTTACGCTACATGACATT 810
                                                                                                                                                                                                                                                              cerrarectricardacarr 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         22-FEB-2000; 2000US-0183756P.
20-OCT-2000; 2000US-00692414.
24-JAN-2001; 2001US-00768184.
                                                                                                                                                                              0.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                20-FEB-2001; 2001WO-US005358.
                                                                                                                                                                                                                                                                                                                                ABA89986 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                      (first entry)
                                                                                                                                                                                                         16; Conservative
 microsatellite markers,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Cassel MJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2002-041152/05.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (PEKE ) PE CORP NY
                                                                                                                                                                                             Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WO200162969-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                      11-FEB-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      30-AUG-2001.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Kalush F,
                                                                                                                                                                                                                                                               7
                                                                                                                                                                                                                                                                                                                                                           ABA89986;
                                                                                                                                                                                Query Match
                                                                                                                                                                                             Local
                                                                                                                                                                                                                                                                                                      RESULT 1030
                                                                                                                                                                                             Best Loca
Matches
                                                                                                                                                                                                                                                                                                                    ABA89986
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mediated/modulated by an oestrogen receptor (ER). (II) is also useful in gene therapy for treating cancer, osteoporosis and cardiovascular diseases. The human ESR-alpha gene is located on chromosome 6. ABA89973 to ABA9010 represent PCR primers, and ABA90011 to ABA90037 represent sequencing primers, for the human ESR-alpha gene, which are used in an example from the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human; calreticulin; antisense compound; hyperproliferative disorder; cancer; autoimmune disease; viral infection; cardiovascular disease; antisense therapy; cytostatic; immunosuppressive; virucide; antisense; phosphorothioate backbone; ss.
                                                                                                                                                                                                                                                   Gaps
                                                                                                                                                                                                                                                   .
                                                                                                                                                                                                    Score 14.2; DB 1; Length 20; Pred. No. 8.7e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human calreticulin antisense oligonucleotide, ISIS 109325.
                                                                                                                                                                                                                                                   3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    note = "2'methoxyethyl nucleotides"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       'note = "2'methoxyethyl nucleotides
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /note= "Phosphorothioate backbone"
                                                                                                                                                           Sequence 20 BP; 2 A; 7 C; 4 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                 Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Location/Qualifiers
                                                                                                                                                                                                                                                                                               826 TCCCTCACCCTTGTCTTG 844
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /mod base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /mod_base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /mod_base= OTHER
                                                                                                                                                                                                                                                   0;
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/mod_base= m5c
6..20
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/mod_base= m5c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /*tag= d
/mod_base= m5c
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/mod_base= m5c
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/mod_base= m5c
                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAD39532 standard; DNA; 20 BP
                                                                                                                                                                                                       0.8%;
                                                                                                                                                                                                                           84.28;
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'*tag= b
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                                                                                                                                                                                                                                                   Conservative
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The invention relates to an antisense compound 8 to 30 nucleotides in
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                  4; Page 33; 528pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (ISIS-) ISIS PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Cowsert LM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2002-583481/62.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WO200248168-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           11-OCT-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Baker BF,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ABT05202;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          RESULT 1033
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                  Claim
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0;
                                                                                                                                                                                                                                                                                                              for modulating the expression of calreticulin. The compositions comprise antisense compounds, particularly antisense oligonucleotides, targetted to nucleic acids encoding calreticulin. The antisense compound is useful for inhibiting the expression of calreticulin in human calls or tissues. It is also useful for treating a human having a disease or condition cancer, autoimmune disease, viral infection or cardiovascular disease, pinhibiting expression of calreticulin. It is useful for diagnostics, therapeutics, prophylaxis and as research reagents and kits. It is also used in antisense therapy. The present sequence is an antisense compound targetted to human calreticulin. This sequence is an antisense compound antisense inhibition of calreticulin expression-phosphorothioate 2'-MOB
                                                                                                                                                                                               Novel antisense compound targeted to nucleic acid encoding calreticulin, useful for treating a human having disease or condition associated with calreticulin e.g. cancer, viral infection, autoimmune disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human; chromosome 1p36-35; chromosome 21q22.1; genetic analysis; genome;
                                                                                                                                                                                                                                                                                                     The invention relates to antisense compounds, compositions and methods
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           .;
0
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84.2%; Pred. No. 8.7e+02;
Live 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human chromosome 1p36-35 PCR primer SEQ ID NO:1451.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Seguence 20 BP; 2 A; 7 C; 6 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                  Claim 3; Page 82; 109pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        928 CAGCTGCTCCGTGGCCTGG 946
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2 CAGCTCGTCCTTGGCCTGG 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ABL44407 standard; DNA; 20 BP.
                                                                 30-OCT-2000; 2000US-00702327.
                                30-OCT-2001; 2001WO-US049045.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               10-MAR-2000; 2000JP-00066716.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                12-MAR-2001; 2001JP-00068285.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (RIKA ) RIKAGAKU KENKYUSHO.
(GENO-) GENOTEX YG.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           11-APR-2002 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                                 Cowsert LM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       gapmer oligonucleotides
                                                                                                (ISIS-) ISIS PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Arraying genome clones.
                                                                                                                                                                  WPI; 2002-479759/51.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2002-144136/19.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              JP2001321190-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             20-NOV-2001.
10-MAY-2002
                                                                                                                                 Bennett CF,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ABL44407;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
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multiwell plates numbered for discrimination are mixed in each of the multiwell plates; (a) a primer designed based on the chromosome marker comultiwell plates; (b) a primer designed based on the chromosome marker comultiwell plates; (b) a primer designed based on the chromosome marker sequence is added to the mixture to carry out an amplification reaction; (c) a signal corresponding to the marker is detected from the resultant amplified product to specify the discrimination Nos. of the multiwell plates containing the clones having said marker sequence; (d) the order of the markers is changed so that the same discrimination Nos. succeed to the marking in the specified discrimination Nos. to array the multiwell plates; (e) the clones in the mixed respectively in each wells of longitudinal and lateral directions; (f) the mixed clones are cultured and the resultant cultures are amplified products; (h) the clones in the multiwell plates are specified from the detected from the detected from the detected from the detected result; and (i) the clones are reconstituted as the positions on the chromosome and arrayed. The mixed respectively is useful for gene analysis. ABL42957 to ABL45322 represent correspondent for human chromosome 21q22.1, which are PERFERENT or human chromosome 21q22.1, which are specifically claimers for human chromosome 21q22.1, which are specifically claimed for use in the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Novel antisense compound targeted to nucleic acid molecule encoding tumor necrosis factor receptor 1 (TNFR1), useful for treating humans having disease associated with TNFR1 e.g. hepatitis, liver injury, liver cancer.
present invention describes a method of arraying genome clones. The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Antisense compound; tumour necrosis factor receptor 1; liver disease; TNFR1; hepatitis; liver injury; hyperproliferative disorder; cancer; mouse; murine; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TNFR1 expression modulation related antisense oligo SEQ ID No 232.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0.8%; Score 14.2; DB 1; Length 20; 34.2%; Pred. No. 8.7e+02; ve 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 20 BP; 5 A; 4 C; 6 G; 5 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1526 TTCAGCTACAAAAGGAGGC 1544
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Zhang H,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1 rrcagcracerardeadec 19
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match 0.8%;
Best Local Similarity 84.2%;
Matches 16; Conservative
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screening of candidate pharmaceutical agents, where high-throughput screening techniques are employed. (II) is useful to detect and quantitate gene expression in biological samples. Oligonucleotides or longer fragments derived from (II) are useful as probes in a microarray used to monitor the expression level of large number of genes. (I)-(III) are useful for the study of the function of a GABA receptor, to study the mechanism of the disease as related to GABA receptor, for the creation of the evaluation of potential therapeutic interventions. ABKZ7339 represent mutant gamma-minobutyric acid receptor subunit coding

886666666666666

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Gaps

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3; Indels

0; Mismatches

ò g

Matches 16; Conservative

Local Similarity

Score 14.2; DB 1; Length 20; Pred. No. 8.7e+02;

0.8%;

Sequence 20 BP; 6 A; 7 C; 4 G; 3 T; 0 U; 0 Other;

sequences and PCR primers of the invention

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length targeted to nucleic acid molecule encoding tumour necrosis factor receptor 1 (TWFR1), where the antisense compound inhibiting expression of TWFR1. The antisense compound is useful for inhibiting the expression of TWFR1 in cells or tissues. The antisense compound is also useful for treating an animal (preferably human) having a disease or condition associated with TWFR1, e.g. a liver disease (such as hepatitis, or liver injury) or a hyperproliferative disorder such as cancer, by inhibiting the expression of TWFR1. The antisense compound is useful for diagnostics, therapeutics, prophylasis and as research reagents and kits. This polynucleotide sequence represents a mouse oligonucleotide relating
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mutant gamma-aminobutyric acid receptor GABARD subunit PCR primer #15
                                                                                                                                                                                                                                                 Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Nootropic; Neuroprotective; Neuroleptic; Antimigraine; Anorectic; adman-aminobutyric acid receptor subunit; GABA; epilepsy; anxiety; manic depression; phobic obsessive symptom; Alzheimer's disease; schizophrenia; migraine; obesity; receptor; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Anticonvulsant; Tranquiliser; Antimanic; Antidepressant;
                                                                                                                                                                                                                                                 .
0
                                                                                                                                                                                                                 0.8%; Score 14.2; DB 1; Length 20;
84.2%; Pred. No. 8.7e+02;
                                                                                                                                                                                                                                                 3; Indels
                                                                                                                                                                                   Sequence 20 BP; 2 A; 6 C; 7 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                 0; Mismatches
                                                                                                                                                                                                                                                                              1565 TGCCTGACTCAGGCAGGCC 1583
                                                                                                                                                                                                                                                                                                       1 TGGCTGGCTCAGGCAGTCC 19
                                                                                                                                                      to the TNFR1 of the invention
                                                                                                                                                                                                                                                                                                                                                                                    ABK27372 standard; DNA; 20 BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              20-JUN-2000; 2000AU-00008260.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             13-SEP-2000; 2000AU-00000098
11-MAY-2001; 2001AU-00004953
                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
                                                                                                                                                                                                                                                 16; Conservative
                                                                                                                                                                                                                                  Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WO200198486-A1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                     ABK27372
                                                                                                                                                                                                                     Query Match
                                                                                                                                                                                                                                                                                                                                                         RESULT 1034
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Fungal; pathogen; banana; polymerase chain reaction; Mycosphaerella; internal transcribed spacer; ITS; PCR primer; ss.

Mycosphaerella sp. WO200196600-A2

Synthetic

20-DEC-2001.

(SYGN) SYNGENTA PARTICIPATIONS AG.

WPI; 2002-130742/17.

Barnett CJ,

16-JUN-2000; 2000US-0211902P. 15-JUN-2001; 2001WO-EP006783.

Mycosphaerella species ribosomal gene-specific primer ITS2.

(first entry)

09-APR-2002

ABA94547;

BP.

ABA94547 standard; DNA; 20

RESULT 1035 ABA94547

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tissue
                                                                                                                                                                                                                                                                                                                                                                                                            Novel oligonucleotide primer useful for polymerase chain reaction-based detection of Mycosphaerella species, a banana fungal pathogen.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                chain reaction (PCR) based detection of a Mycosphaerella sp., a fungal pathogen of bandar. The method involves isolating DNA from a plant tissu infected with Mycosphaerella sp., amplifying a part of ITS (internal transcribed spacer) sequence using the DNA as template in PCR with the specified primer pairs and detecting Mycosphaerella sp. by visualizing the amplified part of ITS sequence. The primers enable the detection of specific isolates of fungal pathogens and the monitoring of disease development in plant populations. Sequences ABA94546-549 represent ribosomal gene-specific for the ITS regions
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The invention relates to oligonucleotide primers for use in polymerase
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DB 1; Length 20; Score 14.2; 0.8%; Query Match

The invention relates to an isolated mammalian polypeptide (I), which is a mutant of gamma aminobutyric acid (GABA) receptor subunit. The mutation distripts the functioning of an assembled GABA receptor, its functional fragment or homologue, and creates a phenotype of epilepsy, anxiety, manic depression, phobic obsessive symptoms, Alzheimer's disease, schizophrenia, migraine and/or obesity. (I), the polymucleotide (I) encoding (I) and antibody (III) to (I) are useful in the diagnosis of epilepsy, anxiety, manic depression, phobic obsessive symptoms. Alzheimer's disease, schizophrenia, migraine and/or obesity. (III) is useful for treating the above conditions. (I)-(III) are useful in

Mutant gamma-aminobutyric acid receptor subunits and DNA molecule, useful for diagnosing epilepsy, Alzheimer's disease, mígraine, obesity, anxiety, manic depression and schizophrenia.

Example 5; Page 52; 99pp; English.

Dibbens LM;

Harkin LA,

Berkovíc SF,

Wallace RH, Mulley JC, (BION-) BIONOMICS LTD.

WPI; 2002-122280/16.

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Ribosome RNA gene base sequence of Cordyceps sinensis for classification of seeds of Cordyceps sinensis.
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                                                                                                      Ribosome ribonucleic acid; rRNA; Cordyceps crassispora; classification;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The invention relates to a novel base sequence which is part of a fully defined ribosome ribonuclaic acid (KRNA) gene of Cordyceps crassispora. The base sequences can be used for the classification of Cordyceps sinensis. The sequence represents a PCR primer used in the invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Disclosure; Page 11; 33pp; Japanese.
                                                                                                                                    Cordyceps sinensis; ss; PCR; primer.
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                                                                                                                                                                                                                                                                                                                                                                                                                                            12-JAN-2001; 2001JP-00004805.
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                                           Cordyceps PCR primer ITS3.
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KANESHIRO N.
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                                                                                                                                                                                                                                                         JP2002204696-A.
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(KANE/)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Novel oligonucleotide primer useful for polymerase chain reaction-based detection of Mycosphaerella species, a banana fungal pathogen.
                                           Gaps
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internal transcribed spacer; ITS; PCR primer; ss.
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llarity 84.2%; Pred. No. 8.7e+02;
Conservative 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mycosphaerella species ribosomal gene-specific primer ITS3.
                                       Indels
      Pred. No. 8.7e+02;
0; Mismatches 3;
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                                                                                             1549 CTTCGGTCTTCGTCGATGC 1567
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                                                                                                                                           2 CTGCGTTCTTCATCGATGC 20
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Best Local Similarity 84.2%;
Matches 16; Conservative
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es 16; Conserv
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ABV78756;

ABV78756/c ID ABV78 XX ABV78 AC ABV78 XX 14-J3

RESULT 1037

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Query Match

Best Loca Matches

Synthetic.

RESULT 1036
ABA94546/C
ID ABA94546/X
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ABA94548

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10-SEP-2002
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             The invention relates to a novel base sequence which is part of a fully defined ribosome ribonucleic acid (rRNA) gene of Cordyceps crassispora. The base sequences can be used for the classification of Cordyceps sinensis. The sequence represents a PCR primer used in the invention
                                                                                                                                                                                                             Human, E2F transcription factor 2, hyperproliferative disorder, cancer, developmental disorder, antisense, therapy, phosphorothioate backbone;
                                                                                 Gaps
                                                                                 ;
0
                                                                                                                                                                                               Human E2F transcription factor 2 antisense oligo, ISIS #114100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /mod_base= OTHER
/note= "2'-methoxyethy1 (2'-MOE) nucleotides"
                                                                                                                                                                                                                                                                                                             *tag= b
mod_base= OTHER
/note= "2'-methoxyethyl (2'-MOE) nucleotides"
                                                                0.8%; Score 14.2; DB 1; Length 20; ilarity 84.2%; Pred. No. 8.7e+02; Conservative 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                               note= "Phosphorothioate backbone"
                                                   Sequence 20 BP; 2 A; 6 C; 5 G; 7 T; 0 U; 0 Other;
 Disclosure; Page 11; 33pp; Japanese.
                                                                                                                                                                                                                                                                  location/Qualifiers
                                                                                               1549 CTTCGGTCTTCGTCGATGC 1567
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es 16; Conserv
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                                                                                                                                                                                 16-JUL-2002
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                                                                                                                                                                   AAD34903;
                                                                    Query Match
                                                                                                                                     RESULT 1039
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Matches
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The present invention relates to antisense oligonuclectides, compounds and methods for modulating the expression of E2F transcription factor 2. The antisense oligonuclectides specifically hybridise with and inhibit the expression of E2F transcription factor 2. They are useful for inhibiting the expression of E2F transcription factor 2 and for treating diseases or conditions associated with E2F transcription factor 2, such as hyperpolliferative disorders, particularly cancer and developmental disorders. They may also be used as research reagents and diagnostics, to distinguish between functions of various members of a biological pathway and in the treatment of a disease or disorder which can be treated by modulating the expression of E2F transcription factor 2. The oligomeric compounds, particularly the antisense oligonucleotides encoding E2F transcription factor 2, ultimately modulating the amount of E2F transcription factor produced. Sequences of the invention are also used in antisense therapy. The present DNA sequence is human E2F transcription factor 2 antisense oligonucleotide with a phosphorothioate backbone. This sequence is targetted to the coding region of human E2F transcription
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New antisense oligonucleotides targeted to a nucleic acid encoding E transcription factor 2, useful for treating a disease or condition associated with E2F transcription factor 2, e.g. hyperproliferative
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Pred. No. 8.7e+02;
0; Mismatches 3; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 20 BP; 1 A; 9 C; 5 G; 5 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 3; Page 92; 120pp; English.
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/mod_base= m5c
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Best Local Similarity 84.2%;
Matches 16; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          disorders, such as cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                 (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Popoff I, Wyatt JR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2002-329864/36.
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The invention describes a compound, 8-50 nucleobases in length targeted to a nucleic acid molecule encoding human TERT (telomerase reverse transcriptase), where the compound specifically hybridises with and inhibits the expression of TERT, a series of oligonucleotides were compound for the human TERT RNA. These were 20 nucleotides in length and composed of a central gap region consisting of the 12-deoxynucleotides, flanked on both sides ($' and 3' directions) by five-nucleotide wings. The wings were composed of 2' methoxyethyl (2' MOB) nucleotides in the compounds were analysed for their effect on human TERT mRNA levels by reverse transcriptase (RT)-polymerase chain reaction (PCR). The compound is useful for inhibiting the expression of TERT in associated with TERT, for modulating apoptosis, for inhibiting cell growth (preferably, cancer cell growth), in antisense therapy and for diagonucleotide used to modulate the activity of nucleic acid molecules encoding TERT, described in the method of the invention
transcriptase (TERT), which specifically hybridizes with and inhibits expression of TERT, useful for modulating apoptosis and inhibiting cell growth.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human, K-ras, PCR primer; probe, capture probe; mutation detection; ligase detection reaction; LDR; p53; BRCA1, BRCA2; infectious disease; infection; 21 hydroxylase deficiency; Turner Syndrome, obesity; cancer; oncogene; tumour suppressor; human papillomavirus; forensic; environmental monitoring; food industry; feed industry; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Designing capture oligonucleotide probes for use on a support to which complementary oligonucleotides hybridize with little mismatch.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0.8%; Score 14.2; DB 1; Length 20; 84.2%; Pred. No. 8.7e+02; tive 0; Mismarches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Seguence 20 BP; 7 A; 9 C; 2 G; 2 T; 0 U; 0 Other;
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                                                                                          Page 91; 154pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (CORR ) CORNELL RES FOUND INC.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ABI95967 standard; DNA; 20
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les 16; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WO200179548-A2.
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                                                                                          Claim 26;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Synthetic
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Matches
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                                                                                                                                                                                                                                                                                                                                                     The present invention relates to a method of minimising immunological transfer (NT) foctus by transferring a nuclear transfer (NT) foctus by transferring a nuclear transfer embryo into an embryo recipient under conditions effective for the development of a nuclear transfer foctus with minimal risk of minumulogical rejection of the foctus due to materanal anti-foctal major histocompatibility complex (MHG)-I immune response. The method is useful for minimising immunological rejection of a NT foctus. It is also useful for parforming embryo transfer. The present DNA sequence is a PCR primer which is used for amplifying bovine MHC class I exon 2 DNA. This sequence is used in the exemplification of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Telomerase reverse transcriptase, TERT, cytostatic, apoptosis, cell growth inhibitor; antisense oligonucleotide, antisense technology;
                                                                                                                                                                                                                           יישות במונים ומשתחוס logical rejection of nuclear transfer fetuses, by transferring the nuclear transfer embryo into an embryo recipient for development of the fetus.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Telomerase reverse transcriptase, antisense oligonucleotide #76.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0.8%; Score 14.2; DB 1; Length 20; 34.2%; Pred. No. 8.7e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0; Mismatches
                                                                                                                                                                                                                                                                                                                      Example 1; Page 71; 103pp; English.
                                                                                                                                                                   Hill JR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1593 CGTGGTGGACACCGAGTTC 1611
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                                                                                                                            (CORR ) CORNELL RES FOUND INC.
                                                   03-OCT-2001; 2001WO-US030925.
                                                                                      03-OCT-2000; 2000US-0237673P.
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07-DEC-2000; 2000US-00733294.
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                                                                                                                                                                 Schlafer DH,
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                                                                                                                                                                                                      WPI; 2002-444101/47.
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es 16; Conserv
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              11-APR-2002
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                                                                                                                                                                   Davies CJ,
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oligonuclectide probes (I) for use on a support to which complementary oligonuclectide probes (II) will hybridise with little mismatch, where clip wave melting temperatures within a narrow range. The method is useful for detecting infectious diseases caused by bacterial infectious agents of a Salmonella, Listeria monocytogenes and Haemophilus influenza, fungal infectious agents e.g. Cryptococcus neoformans, Candida albicans and appergillus funigatus, viruses e.g. T-cell lymphocytotrophis citus, Epstein-Barr virus and pollo virus, and parasitic infectious agents celected from Onchoverva volvulus. Entameba histolytica and Dracunculus medinesis. The method is also useful for detecting genetic diseases such as 1 hydroxylase deficiency, furner Syndrome and obesity defects. Detecting cancer involving oncogenes, tumour suppressor genes or genes involved in DNA amplification, replication, recombination or repair, the nethod is also used for environmental monitoring, forensics and the food and the all passes canning electron microscope and infrared microscope) the support at the particular sites and identifying if ligation of the oligonuclectide probe particular sites and infrared microscope) the support at the particular sites and infrared microscope) the support at the particular sites and infrared microscope) the support at the particular sites and infrared microscope) the support at the particular sites and infrared microscope) the support at the particular sites and infrared microscope) the support at the particular sites and infrared microscope) the support at the particular sites and infrared microscope) the support at the particular sites and infrared microscope) the support at the particular sites and infrared microscope) the support at the particular sites and infrared microscope) the support at the particular sites and infrared microscope and i
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Best Local Similarity 84.2
Matches 16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   the present invention
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The present invention describes a method (M1) for designing capture

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oligonucleotide probes (I) for use on a support to which complementary oligonucleotide probes (II) will hybridise with little mismatch, where clid wave melting temperatures within a narrow range. The method is useful for detecting infectious diseases caused by bacterial infectious agents e.g. Salmonalla, Listeria monocytogenes and Haemophilus influenza, fungal infectious agents and color with a monocytogenes and Haemophilus influenza, fungal specially fundatus, viruses e.g. T-cell lymphocytotrophis citus, Epstein-Barr virus and pollor virus, and parashite infectious agents selected from Onchoverva volvulus. Brianceba histolytica and Dracunculus medinesis. The method is also useful for detecting genetic diseases such so 1 hydroxylase deficiency, Turner Syndrome and obesity defects. Detecting cancer involving oncogenes, tumour suppressor genes, or genes involved in DNA amplification, replication, recombination or repair, the cancer is specifically associated with a gene selected from BRCA1 gene, p53 gene, human papillomavirus types 16 and 18 and liver cancers. The method is also used for environmental monitoring, forensics and the food and for environmental monitoring (using e.g. scanning electron microscope and infrared microscope) the support at the particular sites and identifying if ligation of the oligonucleotide probe presence or absence of the target nucleotide sequences. AB192074 to AB19546 represent oligonucleotide sequences used in the exemplification
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 20 BP; 2 A; 7 C; 5 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Kliman
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0.8%; Score 14.2; DB 1;
84.2%; Pred. No. 8.7e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Capture oligonucleptide Zip ID#235 oligo #9.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Favis R,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example 5; Fig 29; 300pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        999 GCTCATCAACGAGAGGGA 1017
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gerry NP,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (CORR ) CORNELL RES FOUND INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        14-APR-2000; 2000US-0197271P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              19 GCTCATCACAGAGACGGGA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                04-APR-2001; 2001WO-US010958.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ABI93148 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match 0.8
Best Local Similarity 84.2
Matches 16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                            the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2002-034366/04.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WO200179548-A2.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ABI93148;
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oligonuclectide probes (I) for use on a support to which complementary oligonuclectide probes (II) will hybridise with little mismatch, where (I) have melting temperatures within a narrow range. The method is useful for detecting infectious diseases caused by bacterial infectious agents of salmonella, listeria monocytogenes and Heemophilus influenza, fungal infectious agents e.g. Cryptococcus neoformans, Candida albicans and Appergillus funigatus, viruses e.g. T-cell lymphocytotrophis cirus, Epstein-Barr virus and pollo virus, and parasitic infectious agents celected from Onchoverva volvulus, Entamoba histolytica and Dracunculus medinesis. The method is also useful for detecting genetic diseases such as 21 hydroxylase deficiency, Turner Syndrome and obesity defects.

Detecting cancer involving oncogenes, tumour suppressor genes, or genes involved in DNA amplification, replication, recombination or repair, the cancer is specifically associated with a gene selected from BRCA1 gene, p33 gene, human papillomavirus types 16 and 18 and liver cancers. The method is also used for environmental monitoring, forensics and the food and federating comprises scanning quing e.g. as scanning electron microscope and infrared microscope) the support at the particular sites and identifying if ligation of the oligonucleotide probe sets.

MENDALA COLUME and confidention sequences. ABIS2074 to a presence or absence of the target nucleotide sequences. ABIS2074 to a ball of the contraction of the exemplification.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Peptide of estrogen receptor alpha genes variant or its fragment for use in identifying modulators for treating disorders e.g. a susceptibility to cancer, osteoporosis, cardiovascular disorder.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human, oestrogen, receptor, oestrogen receptor alpha, cytostatic,
osteopathic, cardiant, cancer, osteoporosis, cardiovascular disorder,
ESR-alpha, ESR1, PCR, primer, ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ·.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0.8%; Score 14.2; DB 1; Length 20; 84.2%; Pred. No. 8.7e+02; ive 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 20 BP; 6 A; 6 C; 5 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Winn_deen ES;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human ESR1 exon 1G reverse PCR primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1121 TGCTTGGGTCCACGGACTA 1139
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example 1; Fig 2D; 352pp; English
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            recrircedrocaredacea 1
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13-MAR-2001; 2001US-00804076.
05-APR-2001; 2001US-00826314.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ABQ87695 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Local Similarity 84.2
                                                                                                                                                                                                                                                                                                                                                                                                                                             the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Cassel MJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (APPL-) APPLERA CORP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         18-SEP-2002
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            The invention relates to novel human oestrogen receptor variant peptides, and the polymucleotides encoding them. The peptides of the invention have cytostatic, osteopathic and cardiant activity. The peptides of the invention are useful to mediate or modulate a variety of disorders such as a susceptibility to cancer, osteoporosis, cardiovascular disorder, sequences shown in ABQ97682-ABQ9719 represent PCR primers used in the invention to amplify individual exons of the human oestrogen receptor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The invention relates to a novel antisense compound, which is 8-10 calceledies in length targeted to a nucleic acid molecule encoding human calreticulis, and specifically hybridises with and inhibits the expression of human calreticulin. A compound of the invention has expression of human calreticulin. A compound of the invention has and may act as a calreticulin-inhibitor, and have a use in antisense gene therapy. The antisense compound is useful for treating a disease or condition associated with calreticulin e.g. melanoma, viral warts, rubella, schiscosomiasis, congenital heart block or osteoporosis. The present sequence is useful as prophylaxis, research research and diagnostic. The present sequence is used in the exemplification of the invention. The sequence is a phosphorothioate oligonucleotide, having 2'MOS wings and a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New antisense compound that specifically hybridizes with and inhibits the expression of human calreticulin, useful for treating diseases e.g. osteoporosis or schistosomiasis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ss; human; antisense compound; calreticulin; cytostatic; cardiant; virucide; osteopathic; antiparasitic; antisense gene therapy; melanoma; viral warts; rubella; schistosomiasis; congenital heart block;
                                                                                                                                                                                                                                                                                                  Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Phosphorothicate oligonuclectide calreticulin inhibitor SEQ ID NO:66
                                                                                                                                                                                                                                                                                                  .
0
                                                                                                                                                                                                                                                         0.8%; Score 14.2; DB 1; Length 20; 34.2%; Pred. No. 8.7e+02;
                                                                                                                                                                                                                                                                                                Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Bennett CF, Rothlein R, Kishimoto TK, Cowsert LM;
                                                                                                                                                                                                                  Seguence 20 BP; 2 A; 7 C; 4 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Example 15; SEQ ID NO 66; 110pp; English.
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                                                                                                                                                                                                                                                                                                                                         826 TCCCTCACCCTTGTCTTTG 844
                                                                                                                                                                                                                                                                                                                                                                           1 receachecerrereries 19
                                                                                                                                                                             alpha (ESR-alpha or ESR1) gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADG34600 standard; DNA; 20 BP.
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                                                                                                                                                                                                                                                                            84.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
                                                                                                                                                                                                                                                                                                  Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2002-750420/81.
                                                                                                                                                                                                                                                                            Best Local Similarity
Matches 16; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (BOEH ) BOEHRINGER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WO200268688-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             26-FEB-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        osteoporosis
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADG34600;
                                                                                                                                                                                                                                                           Query Match
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deoxy gap.

2001US-0294473P. 2001US-0296964P.

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                                                                                                                                                                                         PCR; ss; NOVX; metabolic disorder; diabetes; anorexia; cancer; cardiovascular; infectious; neurodegenerative; immune; haematopoietic disease; dyslipidaemia; anorectic; virucide; nootropic; antiniflammatory; neuroprotective; antilipaemic; anabolic; cardiant; neurogenesis; wound healing; angiogenesis; chromosome mapping; tissue typing; preventive medicine; pharmacogenomic; primer; human.
                                              Gaps
                                              ö
                                                                                                                                                                           Reverse PCR primer used to amplify human NOVX DNA SeqID1149.
                           Ouery Match

0.8%; Score 14.2; DB 1; Length 20;
Best Local Similarity 84.2%; Pred. No. 8.7e+02;
Matches 16; Conservative 0; Mismatches 3; Indels
            Sequence 20 BP; 2 A; 7 C; 6 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                            07-FEB-2001; 2001US-0266975F.

07-FEB-2001; 2001US-0267459F.

08-FEB-2001; 2001US-0267459F.

08-FEB-2001; 2001US-0267459F.

15-FEB-2001; 2001US-0268974F.

26-FEB-2001; 2001US-0271839F.

27-FEB-2001; 2001US-0271839F.

02-MAR-2001; 2001US-0271858F.

02-MAR-2001; 2001US-0273946F.

14-MAR-2001; 2001US-0273046F.
                                                                 928 CAGCTGCTCGTGGCCTGG 946
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2001US-0288327P.
2001US-0288504P.
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2001US-0275989P.
2001US-0276448P.
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2001US-0276397P.
2001US-0276768P.
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2001US-0278775P.
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2001US-0266767P.
                                                                                                                              BP.
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                                                                                                                                                                                                                                                                                                           31-JAN-2002; 2002WO-US002785
                                                                                                                             ADI17613 standard; DNA; 20
                                                                                                                                                              15-APR-2004 (first entry)
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14-MAR-2001; 2
15-MAR-2001; 2
15-MAR-2001; 2
16-MAR-2001; 2
20-MAR-2001; 2
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27-FEB-2001;
02-MAR-2001;
02-MAR-2001;
14-MAR-2001;
14-MAR-2001;
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05-FEB-2001; 2
07-FEB-2001; 2
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29-MAR-2001;
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31-JAN-2001;
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                                                                                                                                                                                                                                                             Homo sapiens
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                                                                                                                                                 ADI17613;
                                                                                                               RESULT 1047
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This invention relates to a novel nucleic acids, and encoded polypeptides continuously which have properties related to the stimulation of biochemical responses in a cell, tissue, organ or organism.

or physiological responses in a cell, tissue, organ or organism.

continuously, it refers to the use of biologically active fragments for specifically, it refers to the use of biologically active fragments for diagnostic and prognostic assays and furthermore in the treatment of diverse pathological conditions. The present invention describes novel diverse pathological conditions. The present invention describes novel and multiple as methods to modulate their hamped and multiple and eligos, ribozymes and peptide nucleic acids.

Context, cancer, cardionant for treating metabolic disorders, diamente context, cancer, cardiovascular, infectious, neurodespenerative, immune and hammatopoietic diseases as well as various dyslipidaemias.

Context, cancer, cardious have many activities including anorectic, and haematopoietic diseases as well as various dyslipidaemias.

Context, these molecules have many activities including anorectic, contentify small molecules that modulate or inhibit, for example, contenting and cardiant. Furthermore, they are useful in screening assays anabolic and cardiant. Furthermore, they are useful in screening assays anabolic and cardiant. Furthermore, they are useful in corcening act as a context as in chromosome mapping, tissue typing, preventive medicine and cut of pharmacogenomics. This oligonucleotide is a PCR primer used to amplify human NOVX DNA of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New NOVX polypeptides and nucleic acids, useful for preventing or treating NOVX associated disorders, e.g. cancer, cardiomyopathy, atheroscierosis, or diabetes, and in chromosome mapping, tissue typing or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          n BD, Patturajan M, Shimkets RA;
Anderson DW, Rastelli L, Miller CE;
Y, Colman SD, Wolenc AR, Pena CEA;
JP, Lepley DM, Rieger DK, Burgess CE;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
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14.2%; Pred. No. 8.7e+02;
ve 0; Mismatches 3; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Colman SD,
Lepley DM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example 2; SEQ ID NO 1149; 1498pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Tchernev VT, Spytek KA, Zerhusen BD
Li L, Gangolli EA, Padigaru M, And
Gerlach VL, Taupier RJ, Gusev VY, (
Purtak K, Grosse WM, Alsobrook JP,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    506 AGGCTACCTGGAGAAGCT 524
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30-MAY-2001; 2001US-0294473P.

08-JUN-2001; 2001US-0296964P.

18-JUN-2001; 2001US-0299324P.

19-JUN-2001; 2001US-0299324P.

13-AUG-2001; 2001US-0312020P.

16-AUG-2001; 2001US-0312908P.

16-AUG-2001; 2001US-031390P.

21-AUG-2001; 2001US-031390P.

28-AUG-2001; 2001US-031390P.
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                                                                                                                                                                                                                                                                                                                                                                                                          12-SEP-2001; 2001US-0318740P.
19-SEP-2001; 2001US-0323379P.
18-OCT-2001; 2001US-0330245P.
                                                                                                                                                                                                                                                                                  28-AUG-2001; 2001US-0315470P.
31-AUG-2001; 2001US-0316447P.
07-SEP-2001; 2001US-0318115P.
07-SEP-2001; 2001US-0318118P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    18-OCT-2001; 2001US-0330308P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            14-NOV-2001; 2001US-0332701P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (CURA-) CURAGEN CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2002-706998/76.
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Antiarteriosclerotic, cardiant, vasotropic, antiinfective, cytostatic, antiinflammatory; inhibitor, antisense gene therapy, atherosclerosis, short heterodimer partner-1; abnormal; lipid, cholesterol metabolism; cardiovascular disease; infection; inflammation; tumour formation; mouse,

WO2003012033-A2.

13-FEB-2003.

Unidentified.

5

Mouse short heterodimer partner-1 expression oligo SEQ ID No 73.

(first entry)

12-JUN-2003

ABT34198;

ABT34198 standard; DNA; 20 BP.

RESULT 1049

ABT34198,

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The invention relates to an antisense compound that is targeted to a nucleic acid encoding inhibitor.kappa B kinase-gamma, specifically and inhibiting its expression. Compounds of the invention are antisense coligonucleotides comprising at least one modified internucleoside comprising at least one modified internucleoside moiety, which is a P-O-methoxyethyl sugar moiety, or at least one modified nucleobase, which is a 5-O-methoxyethyl sugar moiety, or at least one modified nucleobase, which is a 5-methyloytosine. Preferably, the cutiense oligonucleotide is a chimeric oligonucleotide. The compound of the invention is useful for preparing a composition for treating a chifammatory disorder. The methods are useful for inhibiting the capters of inhibitor-kappa B kinase-gamma in cells or tissues, and creating an animal having a disease or condition associated with creating an animal having a disease or condition associated with crepresent antisense oligonucleotides for the inhibitor-kappa B kinase-gamma. Sequences given in ADA44713 ADA44790 crepresent antisense oligonucleotides for the inhibition of human any and an animal having a disease or condition of human and an animal having a disease or condition associated with a condition of human and an animal having a disease or condition of human and an animal having a disease or condition associated with a condition and animal having a disease or condition of human and an animal having a disease or several and animal having a disease or condition associated with a condition and animal having a disease or condition associated with a condition of human and a sequence given in ADA44713 ADA44790 creating an animal having a disease or condition of human and an animal having a disease or condition and animal having a disease or condition of human and an animal having a disease or condition of human and an animal having a disease or condition of human and animal having a disease or condition and animal having a disease or condition of human and animal having a disease or condi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New compound having sequence targeted to nucleic acid encoding inhibitor-kappa B kinase-gamma, useful for preparing composition for treating e.g., cancer, or inflammatory or autoimmune disorder.
                                                                                             Antisense oligonucleotide; cytostatic; immunosuppressive; antiinflammatory; gene therapy; hyperproliferative disorder; cancer; autoimmune; inflammatory disorder; inhibitor-kappa B kinase-gamma; ss;
                                                                                                                                                                                                                                                                                                                                                                  /note= "Phosphorothioate linkages, all cytosines are
methylcytosine"
                                                                                                                                                                                                                                                                                                                                                                                                                                     /*tag= a
/mod_base= OTHER
/note= "2'-methoxyethyl (2'-MOE) nucleotides"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               'note= "2'-methoxyethyl (2'-MOE) nucleotides"
                                                 Antisense oligonucleotide #ISIS 115460 #SEQ ID 86.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 20 BP; 3 A; 8 C; 8 G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                            Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                               mod_base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /*tag= c
/mod_base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 3; Page 78; 106pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    03-OCT-2002; 2002WO-US031809.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   06-OCT-2001; 2001US-00972607.
20-NOV-2003 (first entry)
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                                                                                                                                                                                                                                                                                                                    /*tag=
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                                                                                                                                                                                                                                                                  Key
modified_base
                                                                                                                                                                                                                   Homo sapiens.
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The invention relates to a novel compound of 8 - 50 nucleobases in length targeted to a nucleic acid molecule encoding a short heterodimer partner—1. The novel compound specifically hybridizes with a nucleic acid expression of the nucleic acid expression of the nucleic acid molecule. The compound, and a composition comprising it are useful for treating a disease or condition associated abnormal lipid or cholesterol metabolism such as atherosciercism or addition associated abnormal lipid or cholesterol metabolism such as atherosciercism or addition involving cardiovascular disease. They are also useful in research and diagnostics for modulating the expression of short heterodimer partner—1. They can also be useful prophylactically in preventing or delaying infection, and the numeral manner of the contraction of the numeral manner of the numeral manner of the numeral manner of the numeral manner of the numeral numeral manner of the numeral numera
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  inflammation or tumour formation. This polynucleotide sequence represents a mouse antisense oligo relating to the heterodimer partner-1 of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New antisense oligonucleotide targeted to a nucleic acid encoding short beterodimer partner-1, useful for treating diseases involving abnormal lipid or cholesterol metabolism, e.g atherosclerosis or cardiovascular
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
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Pred. No. 8.7e+02;
0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 20 BP; 9 A; 4 C; 7 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             17-JUL-2002; 2002WO-US023245.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             31-JUL-2001; 2001US-00919197.
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16; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Graham MJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ACC49703 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2003-248161/24.
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ID ACC497
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AC ACC497
XX
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Gaps

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0; Mismatches

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0.8%; Score 14.2; DB 1; Length 20; 84.2%; Pred. No. 8.7e+02;

84.2%;

Query Match Best Local Similarity Matches 16; Conserv

Conservative

366 GAGTGACCAGGCTTCAGCC 384

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GAGAGCCCAGCTTCAGCC

19

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The present invention describes a compound 8-50 nucleobases in length targeted to, and which specifically hybridises with a nucleic acid molecule encoding Kinses suppressor of ras-1 (KiSR), and inhibits the molecule encoding Kinses in length that specifically hybridises with at least an 8-nucleobase in length that specifically hybridises with at least an 8-nucleobase portion of an active site on a nucleocate acid molecule encoding KiR; (2) a composition comprising the compound and a carrier or diluent; (3) inhibiting the expression of KSR in cells or tissues by contacting the compound so that expression of KSR is inhibited; and (4) treating an animal having a disease or condition associated with KSR by administering to the animal a therapeutic or prophylactic amount of the compound so that expression of KSR is inhibitor, and in antisense gene therapy. The compound, composition and methods are useful for treating a disease or condition associated with KSR, such a sising from aberrant apoptosis by inhibiting the expression of KSR. They are also useful in research and diagnostics for modulating the expression of KSR. The present sequence represents a chimeric phosphorothioate from the compound of human KSR, which is used in an example from
                                                                     Human, kinase suppressor of ras-1; KSR; cytostatic; KSR inhibitor; antisense gene therapy; hyperproliferative disorder; phosphorothioate; developmental disorder; antisense oligonucleotide; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New compounds, particularly antisense oligonucleotides targeted to a nucleic acid encoding KSR, useful for treating a disease/condition associated with KSR, such as hyperproliferative or developmental disorders.
                                     Human KSR chimeric phosphorothioate oligonucleotide SEQ ID NO:73.
                                                                                                                                                                                                                                                                                                                                                                                                                          'not\overline{e} = "2' - 0 - methoxyethyls (2' - MOE)"
                                                                                                                                                                                                                                                                                               *tag= b /mod_base= OTHER /mod_base= OTHER (2'-MOE)'
frote= "2'-O-methoxyethyls (2'-MOE)'
frag= c /*tag= c
                                                                                                                                                                                                                                                                         note= "phosphorothioate backbone"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 20 BP; 2 A; 6 C; 7 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 15; Page 75; 102pp; English.
                                                                                                                                                                                                    Location/Qualifiers
                                                                                                                                                                                                                  1..20
/*tag= a
/mod_base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                        mod base= OTHER
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         20-SEP-2001; 2001US-00961001
01-JUL-2003 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (ISIS-) ISIS PHARM INC.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                WO2003025144-A2
                                                                                                                                                                                                    Key
modified_base
                                                                                                                                            Homo sapiens.
Synthetic.
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                                                                                                                                                                                                                                                                                             modified base
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This invention relates to the detection of a fungal pathogen comprising isolating DNA from a plant leaf infected with a pathogen. The methods and primers are useful for identifying fungal isolates of fungal pathogens and monitoring of disease development in plant populations. The present sequence represents an oligonucleotide primer used to detect Fusarium ear
                                                                                                                                                                                                                                                                                                                                                                     Detecting a fungal pathogen, useful for monitoring disease development, comprises subjecting the DNA to PCR amplification using at least one primer having sequence identity with at least 10 contiguous nucleotides
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
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0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0.8%; Score 14.2; DB 1; Length 20; 84.2%; Pred. No. 8.7e+02; tive 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 20 BP; 7 A; 5 C; 6 G; 2 T; 0 U; 0 Other;
                                                                                                                                Mitochondria; fungal pathogen; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mitochondria; fungal pathogen; PCR; primer; ss.
                                                                                                                                                                                                                                                                                          (SYGN ) SYNGENTA PARTICIPATIONS AG.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1549 CTTCGGTCTTCGTCGATGC 1567
                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 6; Page 17; 44pp; English.
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                         BP
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                                                                                                       Oligonucleotide primer ITS3
                        ACC50005 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ACC50004 standard; DNA; 20
                                                                          (first entry)
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nes 16; Conservative
                                                                                                                                                                                                                                                                                                                   Beck JJ, Barnett CJ;
                                                                                                                                                                                                                                                                                                                                            WPI; 2003-363229/34.
                                                                                                                                                                                    WO2003027635-A2
                                                                                                                                                                                                                                                                                                                                                                                                               of Fusarium spp
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    cot pathogens
                                                                           14-JUL-2003
                                                                                                                                                                                                              03-APR-2003
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                                                                                                                                                          Synthetic
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                                                   ACC50005;
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Gaps

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0.8%; Score 14.2; DB 1; Length 20; 84.2%; Pred. No. 8.7e+02; tive 0; Mismatches 3; Indels

Query Match
Best Local Similarity 84.2
Matches 16; Conservative

0;

Gaps

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thermophilus (ABV99898). The plasmid is useful for making a transformant which is used for the preparation of foods, food additives, feeds, research reagents or drugs. The present sequence is a PCR primer, which was used in an example from the invention
                                                                                                                                                                                                                                                                                                                                                                                                Mouse; src-c; tyrosine kinase; src-c inhibitor; cytostatic; osteopathic; antisinfammatory; antibacterial; antisense therapy; vaccine; cancer; antisense oligonucleotide; aberrant bone remodeling; breast cancer; hyperproliferative disorder; pancreatic cancer; lung cancer; tumour; ovarian cancer; oesophageal cancer; neuroblastoma; retinoblastoma; kaposi's sarcoma; infection; inflammation; tumour formation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New antisense oligonucleotides targeted to nucleic acids encoding src-c, useful for diagnosing, treating or preventing diseases associated with the expression of src-c, e.g. cancer or inflammation, and in research
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The present invention describes a compound (I) that is 8-50 nucleobases
                                                                                                                                                                                                                                                                                                                                                                    Mouse src-c chimeric phosphorothioate oligonucleotide SEQ ID NO:147
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (2'-MOE wing)"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /note= "2'-0-methoxyethyl gapmer (2'-MOE wing)"
                                                                                                          Length 20;
                                                                                                                                       Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 'note= "phosphorothioate linkages"
                                                                             Sequence 20 BP; 6 A; 5 C; 6 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /note= "2'-0-methoxyethyl gapmer
16. .20
                                                                                                        Score 14.2; DB 1;
Pred. No. 8.7e+02;
                                                                                                                                       0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 3; Page 92; 137pp; English.
                                                                                                                                                                    208 GAGCAGATAGGCCTGGATG 226
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /mod_base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                'mod_base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             OTHER
                                                                                                                                                                                                   19
                                                                                                                                                                                                                                                                              BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2002WO-US015684.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                18-MAY-2001; 2001US-00860473.
                                                                                                                                                                                                 GAGCATATAGCCCTGGAAG
                                                                                                          0.8%;
                                                                                                                                                                                                                                                                            ABZ59526 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /*tag≈ c
/mod base=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Д
                                                                                                                                                                                                                                                                                                                                         (first entry)
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                                                                                                                                       16; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2003-120806/11.
                                                                                                                        Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WO200295053-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    modified base
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mus musculus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  16-MAY-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    applications
                                                                                                                                                                                                                                                                                                                                         17-APR-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Synthetic
                                                                                                                                                                                                                                                                                                           ABZ59526;
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                                                                                                           Query Match
                                                                                                                        Best Local
Matches
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0
                                                                                                                                                                                                                                                                                                                       This invention relates to the detection of a fungal pathogen comprising isolating DNA from a plant leaf infected with a pathogen. The methods and primers are useful for identifying fungal isolates of fungal pathogens and monitoring of disease development in plant populations. The present sequence represents an oligonucleotide primer used to detect Fusarium ear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The present invention relates to plasmid pMT1 derived from Streptococcus
                                                                                                                                                                                                                Detecting a fungal pathogen, useful for monitoring disease development, comprises subjecting the DNA to PCR amplification using at least one primer having sequence identity with at least 10 contiguous nucleotides of Pusarium spp.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          research reagent; drug; PCR; primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A new plasmid of Streptococcus thermophilus and its derivatives, make a transformant, a food, a food additive, a feed, a research
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ô
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Streptococcus thermophilus plasmid pMT1-related PCR primer #7.
                                                                                                                                                                                                                                                                                                                                                                                                                                                               0.8%; Score 14.2; DB 1; Length 20; 84.2%; Pred. No. 8.7e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0; Mismatches
                                                                                                                       (SYGN ) SYNGENTA PARTICIPATIONS AG.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          food; food additive;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example 3; Page 19; 25pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1549 CTTCGGTCTTCGTCGATGC 1567
                                                                                                                                                                                                                                                                                           Claim 6; Page 17; 44pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (MEIP ) MEIJI MILK PROD CO LTD.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   BP
                                                             19-SEP-2002; 2002WO-US030311
                                                                                           24-SEP-2001; 2001US-00961755
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               02-MAR-2001; 2001JP-00059196
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2001JP-00059196
                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 20 BP; 2 A; 6 C; 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Streptococcus thermophilus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                                        Barnett CJ;
                                                                                                                                                                                  WPI; 2003-363229/34.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2003-096538/09
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ABV99905 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Local Similarity
hes 16; Conserv
WO2003027635-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   JP2002253260-A.
                                                                                                                                                                                                                                                                                                                                                                                                     rot pathogens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Plasmid pMT1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             02-MAR-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               21-FEB-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   10-SEP-2002.
                                 03-APR-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  and a drug.
                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
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                                                                                                                                                      Beck JJ,
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Matches
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coding region, intron region, exon region, stop codon, intron:exon region, exon region, exon region, exon region, exon region, exon region, or stop codon, intron:exon junction, or stop reactivities and which conservation intron and inhibits the expression of Bro-c. (I) specifically hybridises with and inhibits the expression of Bro-c. (I) have cytostatic, antiinflammatory, osteopathic and antibacterial cartivities, and can be used for modulating the expression of src-cand for treating diseases or conditions associated with expression of cand for treating diseases or conditions associated with expression of src-c, e.g. aberrant bone remodeling or hyperproliferative disorders, or and for treating or bycerproliferative disorders, cancer, owarian cancer, oesophageal cancer, neuroblastoma, retinoblastoma cancer, ovarian cancer, oesophageal cancer, neuroblastoma, retinoblastoma cancer, ovarian cancer, osophageal cancer, neuroblastoma, retinoblastoma cancer, response to delay infection, inflammation or tumour prophylaxis, e.g. to prevent or delay infection, inflammation or tumour companies of various members of a biological pathway. The present companies of various members of a biological pathway. The present continoned continoned continoned in an example from the present invention continoned in an example from the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ss; rat; Jun N-terminal kinase; JNK1; JNK2; JNK3; antisense; cytostatic; antiinflammatory; apoptosis; prostate cancer; prostate tumour; inflammation; fibrosis; fibrotic disease; fibrotic scarring; peritoneal adhesion; lung fibrosis; conjunctival scarring; hyperproliferative disease; cancer; probe.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /note= "2'methoxyethoxy-modified and phosphorothioate
linkages"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /mod_base= OTHER
/note= "2'methoxyethoxy-modified and phosphorothioate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Rat Jun N-terminal kinase, JNK1, antisense oligonucleotide ISIS21867
in length targeted to a nucleic acid molecule encoding a 5'UTR, 3'UTR,
                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
                                                                                                                                                                                                                                                                                                                                                                                                         ..
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      note= "All cytosines are 5-methyl-cytosines"
                                                                                                                                                                                                                                                                                                                                                                 0.8%; Score 14.2; DB 1; Length 20;
34.2%; Pred. No. 8.7e+02;
ive 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                  Sequence 20 BP; 2 A; 5 C; 8 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                  1610 TCTAAGCCACAGACCGAGG 1628
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /mod_base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                97US-00910629.
98US-00130616.
99US-00287796.
99US-00396902.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 20 TCCAAGCCTCAGACCCAGG 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            31-JAN-2001; 2001US-00774809.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ВР
                                                                                                                                                                                                                                                                                                                                                                                             84.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADA26668 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                L. .20
/*tag= b
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         *tag= a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     linkages"
                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity 84.28
Marches 16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 16. .20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /*tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Rattus norvegicus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           US2003004120-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   modified base
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        13-AUG-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        07-APR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       07-AUG-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               02-JAN-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADA26668;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               RESULT 1055
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The invention relates to an oligonucleotide (antisense, AS) comprising 8-
30 nucleotides connected by covalent linkages, where the oligonucleotide
30 nucleotides connected by covalent linkages, where the oligonucleotide
30 nucleotides connected by protein and andulates the expression of the
30 Jun N-terminal kinase (JNK) protein and andulates the expression of the
30 Jun N-terminal kinase (JNK) protein and andulates on comprising
31 carrier), treating an animal having/suspected of having/prone to having a
32 carrier), treating an animal having/suspected of having/prone to having a
33 carrier), treating an animal having/suspected of having/prone to having
34 carrier), treating an animal having/suspected of having/prone to having
35 carrier), treating an animal having/suspected of having/prone to having
36 carrier, therapeutic amount of the composition of the AS oligonucleotide,
37 contacting the expression of a JNK protein of the AS oligonucleotide,
38 contacting the expression of a JNK protein, or expression of a protein that
38 prompted by a JNK protein, or expression of a cellular protein that
39 condition associated by a LNK protein, or expression of a cellular protein of a naminal by administering the oligonucleotide to the cells or the cells of the contacting a propholation and treating a human having a disease or condition associated with a JNK protein or characterised by a reduction in apoptosis, such as a classes or condition (e.g. tibrocis carring, peritoneal adhesions, lung disease or condition (e.g. tibrocis carring, peritoneal adhesions, lung tibrosis or conjunctival scarring), hyperproliferative disease or condition (e.g. tibrocis carring, peritoneal adhesions, lung tibrosis or conjunctival scarring), hyperproliferative disease or condition (e.g. tibrocis carring, peritoneal adhesions, lung tibrosis or condition (e.g. tibrocis carring, peritoneal adhesions, lung scarring), hyperproliferative disease or condition (e.g. tibrocis carring, peritoneal adhesions, lung scarring), hyperprolifer mucleic aci
                                                                                                                                                                                                      New oligonucleotides which hybridizes to, and modulates the expression of Jun N-terminal kinase, useful for treating a disease or condition characterized by a reduction in apoptosis, e.g. prostate cancer,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Antisense; interferon gamma receptor 2; autoimmune disorder; cancer; autoimmune thyroiditis; autoimmune insulinitis; multiple sclerosis; diabetes; autoimmune arthritis; Crohn's disease; apoptosis; IFNGR2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human IFNGR2 antisense oligonucleotide, ISIS #142777.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 20 BP; 4 A; 6 C; 5 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0.8%; Score 14.2; DB 1;
34.2%; Pred. No. 8.7e+02;
ve. 0; Mismatches 3;
                                                                                                                               Gaarde WA;
                                                                                                                                    Nero P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1424 GGATCTCCGCAGAGGATGC 1442
                                                                                                                                                                                                                                                                                                                           Example 7; Page 33; 69pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          7
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hes 16; Conservative
                                                                                                                                    Mckay R, Dean NM, Monia BP,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (first entry)
                                                                                                                                                                                                                                                                                   inflammation or fibrosis.
                                                                                                                                                                              WPI; 2003-311908/30.
             MCKAY R.
DEAN N M.
MONIA B P.
                                                                                              (GAAR/) GAARDE W A.
                                                                               NERO P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        02-MAY-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RESULT 1056
               (MCKA/)
(DEAN/)
(MONI/)
                                                                               NERO/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Matches
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/note= "Phosphorothioate backbone; All cytidine residues

are 5-methylcytidines"

/*tag= b

/mod_base= OTHER

...20 *tag=

Location/Qualifiers

(2'-MOE) nucleotides"

/mod_base= OTHER /note= "2'-methoxyethyl 16..20

/*tag= c /mod_base= OTHER

/note= "2

Wyatt JR;

-methoxyethyl (2'-MOE) nucleotides"

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Human; antisense; fibroblast growth factor receptor 3; prophylaxis;
developmental disorder; hyperproliferative disorder; antisense therapy;
FGFR-3; ACH; JTK4; CEK2; cancer; phosphorothioate; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                             Novel compound targeted to a nucleic acid molecule encoding fibroblast growth factor receptor 3, useful for inhibiting the expression of the receptor and for treating an animal having cancer or developmental
                                                                                                                                                                                                                                                                                                                  06-SEP-2002; 2002WO-US028549.
                                                                                                                                                                                                                                                                                                                                       10-SEP-2001; 2001US-00953047.
                                                                                                                                                                                                                                                                                                                                                             (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2003-313244/30.
                                                                                                                                                                                                                                                                      WO2003023004-A2
                                                                                                     Key
modified_base
                                                                                                                                                                      modified base
                                                                                                                                                                                                                modified_base
                                                                      sapiens
                                                                                                                                                                                                                                                                                             20-MAR-2003
                                                                                Synthetic
                                                                                                                                                                                                                                                                                                                                                                                     Monia BP,
                                                                     Homo
             The invention relates to antisense compounds, composition and methods for modulating the expression of human interferon gamma receptor 2 (IFNGR2). The compositions comprise antisense compounds targetted to nucleic acids treating diseases or conditions associated with IFNGR2, e.g. autoimmune thyroiditis, diabetes, multiple sclerosis, autoimmune atthritis, autoimmune insulinitis or Crohn's disease), cancer, for a disease/disorder caused by aberrant apoptosis. They are also useful kits. The invention is useful in gene therapy. The present sequence is an antisense oligonucleotide targetted to human IFNGR2 DNA
                                                                                                     /note= "Phosphorothioate backbone; All cytidine residues are 5-methylcytidines"
                                                                                                                                                                                                                                                                                                                                                                                New antisense oligonucleotides for modulating Interferon gamma receptor 2, particularly useful for treating autoimmune disorders (e.g. multiple sclerosis or Crohn's disease), cancers or diseases caused by aberrant apoptosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0.8%; Score 14.2; DB 1; Length 20; 4.2%; Pred. No. 8.7e+02;
 gene therapy; prophylaxis; human; phosphorothioate; ss.
                                                                                                                                                                                                    /note= "2'-methoxyethyl nucleotides"
                                                                                                                                                            nucleotides'
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 20 BP; 4 A; 3 C; 8 G; 5 T; 0 U; 0 Other;
                                                                                                                                                          note= "2'-methoxyethyl
                                                          Location/Qualifiers
1. .20
                                                                                         mod_base= OTHER
                                                                                                                                              'mod_base= OTHER
                                                                                                                                                                                             OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 3; Page 85; 127pp; English.
                                                                                                                                                                                                                                                                                          26-APR-2001; 2001US-00843377.
                                                                                                                                                                                                                                                                    16-APR-2002; 2002WO-US012007.
                                                                                                                                                                               /*tag= c
/mod_base= (
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nes 16; Conservative
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                                                                                                                                                                                                                                                                                                                (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                                                                                                     Bennett CF, Watt AT;
                                                                                                                                                                                                                                                                                                                                                           WPI; 2003-156688/15.
                                                       Key
modified_base
                                                                                                                                                                                                                        WO200288163-A1
                                                                                                                      modified_base
                                                                                                                                                                 modified base
                      Homo sapiens
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                                  Synthetic
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Matches
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The invention relates to antisense compounds targetted to a nucleic acid molecule encoding fibroblast growth factor (FGF) receptor 3 (also known as FGFR-3, ACH, JTK4 and CEK2) to inhibit its expression. Antisense compounds of the invention are useful for treating diseases or conditions hyperproliferative disorders, such as developmental disorders or hyperproliferative disorders, sepecially cancer of colorectal, bladder, bone, lung, cervical, breast or skin. They are useful as research reagents, therapeutics, prophylaxis, kits and diagnostics, and as tools in differential and/or combinatorial analyses to elucidate expression
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                .;
0
                                                                                                                                                                                                                                                                                                             patterns of a portion of the genes expressed within cells and tissues. They are also useful in antisense therapy. The present sequence is an antisense oligonucleotide targetted to human FGFR-3
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                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 20 BP; 3 A; 8 C; 3 G; 6 T; 0 U; 0 Other;
Claim 3; Page 79; 120pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                335 ACGAGGACTTGAAGATGGG 353
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         20 ACGGGTACCTGAAGATGGG 2
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les 16; Conservative
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Gaps

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Mismatches

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TGCTGAAGCTCAGTGGAGG TGCTGAAACCCAGGGGAGG

62

à g Human FGFR-3 antisense oligonucleotide, ISIS #125204.

(first entry)

07-AUG-2003

AAD55498;

EX PX XX B

AAD55498 standard; DNA; 20 BP.

RESULT 1057

AAD55498,

29-JUL-2003 (first entry)

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The invention relates to a novel fermentation process to be used in the production of exocellular pravastatin and lovastatin which comprises cultivating microorganisms from Aspergillus and Monascus species. Statins cultivating microorganisms from Aspergillus and Monascus species. Statins are fungal secondary metabolites which inhibit hydroxy-methylglutaryl coenzyme A (Hand-CoA) reductase, the first committed enzyme of cholesterol coenzyme A fland-CoA) reductase, the first committed enzyme of cholesterol synthesis. Statins are therefore used as cholesterol-lowering drugs. The gravastatin, either in a cell-associated form or releasable into the pravastatin, directly, as a secondary metabolite, in the fermentation culture medium. Those production processes currently in existence culture medium. Those production which wise the process culture fermentation of complex carbon monascus ruber. In addition, the process uses simple and complex carbon sources obtained from agricultural waste thereby reducing production costs. The current sequence is that of the fungal universal ITS3 PCR primer of the invention which was used to amplify the Aspergillus terreus
                                                                        Fungal; ITS3; interspace 3 region; ss; fermentation process; lovastatin;
exocellular pravastatin production; statin; HMG-CoA; primer; PCR;
cholesterol synthesis; cholesterol-lowering drug;
hydroxy-methylglutaryl coenzyme A reductase.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Fermentation useful for producing pravastatin involves pre-fermenting fungal strain in first nutrient medium, and then fermenting strain in second nutrient medium.
                                     Fungal universal ITS3 PCR primer - used to amplify ITS2 region DNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PCR primer #14 for human oestrogen receptor alpha (ESR1) gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human; oestrogen receptor alpha; ESR1; cancer; osteoporosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 20 BP; 7 A; 5 C; 6 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match 0.8%; Score 14.2; D Best Local Similarity 84.2%; Pred. No. 8.7e Matches 16; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Disclosure, Page 10; 15pp; English.
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                                                                                                                                                                                                                                                                                                                                                  15-JUN-2001; 2001EP-00114462.
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                                                                                                                                                                                                                                                                                                                                                                                                                                     Manzoni M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2003-423103/40.
                                                                                                                                                                                                                                                                                                                                                                                              (GNOS-) GNOSIS SRL
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                                                                                                                                                                                                                                    EP1266967-A1
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Rollini M;

Nichele M,

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a sample, methods of determining a risk of having or developing a disorder mediated by a variant oestrogen receptor and methods for screening compounds useful for treating such disorders. ABX33718-ABX3375s represent PCR primers for the human BSR1 gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The present invention relates to the sequencing of genomic DNA encoding human human oestrogen receptor alpha (ESR1) protein. The gene encoding human ESR1 is located on chromosome 6. The invention provides the genomic structure of the ESR1 gene and novel single nucleotide polymorphisms (SRPs) Maplotypes in the genes. The polymorphisms/haphotypes can lead to a variety of disorders (such as cancer, osteoporosis, and cardiovascular disorders) that are mediated by a variant oestrogen receptor. The invention provides methods of detecting ESR1 polymorphisms/haplotypes in invention provides methods of detecting ESR1 polymorphisms/haplotypes in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Nucleotide sequence of 5'-biotin-labeled universal capture probe ITS3-B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Dimorphic fungus; internal transcribed spacer-2; ITS2; fungal infection;
                                                                                                                                                                                                                                                                                                                                                                                           Novel isolated estrogen receptor alpha variant peptide, useful in development of diagnostics and therapies for diseases or disorders mediated/modulated by the estrogen receptor, or as immunogens to raise
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
cardiovascular disorder; variant oestrogen receptor; ESR1 haplotype; ESR1 polymorphism detection; cytostatic; osteopathic; cardiant; PCR;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0.8%; Score 14.2; DB 1; Length 20; 84.2%; Pred. No. 8.7e+02; ative 0; Mismatches 3; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 1; Fig 2d; 186pp; English.
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                                                                                                                                                                                                          22-FEB-2000; 2000US-0183756P.
20-OCT-2000; 2000US-00692414.
24-JAN-2001; 2001US-00768184.
13-MAR-2001; 2001US-00804076.
05-APR-2001; 2001US-00805314.
                                                                                                                                                                  21-AUG-2001; 2001US-00933267.
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Dest Local Similarity 84.2
Matches 16; Conservative
                                                                                                                                                                                                                                                                                                                                         Kalush F, Cassel MJ,
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                                                                                                                                                                                                                                                                                                             (PEKE ) PE CORP NY.
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                                                                                                           JS2002123095-A1
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                                                                             Homo sapiens
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                                                  primer; ss.
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Indels

Score 14.2; DB 1; Length 20; Pred. No. 8.7e+02;

03-APR-2003

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modified_base
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                                                                                                                                                                                                                                                                                                                                                                                                                  sample, where the presence of the ITS2 nucleic acid sequence indicates the sample was contacted by the dimorphic fungus. The method is useful for detecting or diagnosing fungal infections. The array is useful screening a sample for the presence of, or contamination by a dimorphic fungus. The present sequence represents a 5' biotin-labeled universal capture probe, used for detecting a dimorphic fungus
                                                                                                                                                                                                                                Detecting a dimorphic fungus, useful for diagnosing fungal infections, comprises detecting the presence or absence of an internal transcribed spacer-2 (ITS2) nucleic acid sequence of a dimorphic fungus within a
                                                                                                                                                                                                                                                                                                                                                              The invention relates to detecting a dimorphic fungus. The method involves detecting the presence or absence of an internal transcribed spacer-2 (ITS2) nucleic acid sequence of a dimorphic fungus within a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            "mod_base= OTHER
'note= "Phosphorothioate backbone; All cytidines are 5-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human ABC transporter MHC I antisense oligonucleotide, ISIS 206637.
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84.2%; Pred. No. 8.7e+02;
iive 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 20 BP; 7 A; 5 C; 6 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             'note= "2'methoxyethyl nucleotides"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /note= "2'methoxyethyl nucleotides"
                                                                                                                                                                  Morrison CJ;
                                                                                                                            SERVICES.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1549 CTTCGGTCTTCGTCGATGC 1567
                                                                                                                                                                                                                                                                                                                             Claim 5; Page 35; 71pp; English.
                                                                                                               (USSH ) US DEPT HEALTH & HUMAN
                                                                                                                                                              Choi JS,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /*tag= c
/mod_base= OTHER
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                                       25-SEP-2002; 2002WO-US030605.
                                                                                   2001US-0325241P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     BP
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*tag=
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                                                                                                                                                                                              WPI; 2003-354661/33.
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Best Local Similarity
                                                                           26-SEP-2001;
                                                                                                                                                      Lindsley MD,
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Matches
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where the compound specifically hybridises with the nucleic acid molecule and inhibites expression of ATM or specifically hybridises with the nucleic acid molecule portion of an active site on the nucleic acid molecule. The invention is useful for inhibiting the expression of ATM in cells or tissues. The invention is useful for treating an animal with hyperproliferative or therapeutics, prophylaxis, as research reagents and kits, for therapeutics, prophylaxis, as research reagents and kits, for distinguishing functions of various members of a biological pathway and e.g., to prevent or delay infection, inflammation or tumour formation. The present sequence is an antisense oligo targetted to human ABC transporter MHC I DNA. This sequence is used to illustrate the method of
                                                                                                                                                                                                                                                                                New antisense compound that hybridizes and inhibits the nucleic acid encoding ABC transporter major histocompatibility complex 1, for treating diseases or conditions such as a hyperproliferative or autoimmune
                                                                                                                                                                                                                                                                                                                                                                                                                                                          The invention relates to a compound targetted to a nucleic acid molecule
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1..20
/*tag= a
/mod_base= OTHER
/note= "Phosphorothioate backbone; All cytidine residues
are 5-methylcytidines"
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MyD88; Alzheimer's disease; neurodegenerative disease; schizophrenia;
gene therapy; Down's syndrome; phosphorothioate; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0.8%; Score 14.2; DB 1; Length 20;
84.2%; Pred. No. 8.78+02;
ive 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human MyD88 antisense oligonucleotide, ISIS #190957.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 20 BP; 3 A; 8 C; 6 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                Freier SM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 3; Page 81; 112pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1461 CCTCAGTCTGGGGGGGGGG 1479
12-DEC-2002; 2002WO-US040101.
                                                  17-DEC-2001; 2001US-00024369.
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                                                                                                                                                                 Borchers AH, Ward DT,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                              (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                WPI; 2003-577305/54.
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Best Local Similarity
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WPI; 2003-810907/76.
      WPI; 2003-523102/49.
                                                                                                                                                            Query Match
Best Local Similarity
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modified base
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      19-JUN-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Graham MJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                     modified
                                                                                                                                                                                                                                                                              ADC35560;
                                                                                                                                                                                                                                           RESULT 1064
                                                                                                                                                                     Best Loca
Matches
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                                                                                                                                                                                                                                          The invention relates to antisense compounds targetted to a nucleic acid encoding human MyD88 (myeloid differentiation primary response gene 88) to inhibits its expression. Antisense compounds of the invention are useful for preparing a composition for treating neurodegenerative disease e.g. Alzheimer's disease, Down's syndrome or schizophrenia. The invention is also useful in gene therapy. The present sequence is an antisense oligonucleotide targetted to human MyD88 DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           controller metabolism; weed; herbicide; herbicide-resistant plant;
                                                                                                                                                                                        New antisense compound, having a sequence targeted to a nucleic acid encoding MyD88, useful for preparing a composition for treating neurodegenerative disease, e.g. Alzheimer's disease.
                                                                                                                                                                                                                                                                                                                                                        Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Weed controller metabolism associated PCR primer SEQ ID NO:83.
                                     /mod_base= OTHER
/note= "2'-methoxyethyl (2'-MOE) nucleotides"
           nucleotides"
                                                                                                                                                                                                                                                                                                                                    0.8%; Score 14.2; DB 1; Length 20;
84.2%; Pred. No. 8.7e+02;
tive 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                    Sequence 20 BP; 7 A; 6 C; 2 G; 5 T; 0 U; 0 Other;
  /mod_base= OTHER
/note= "2'-methoxyethyl (2'-MOE)
16. .20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Nakajima H, Mukumoto F, Takaishi M;
                                                                                                                                                                                                                                                                                                                                                                            TTGTCTTTGAGTACCTGGA 854
                                                                                                                                                                                                                               Claim 3; Page 76; 106pp; English
                                                                                                                                                                                                                                                                                                                                                                                              TGGACTTTGAGTACTTGGA 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          agrochemical; ss; PCR; primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                          ADC36216 standard; DNA; 20 BP
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07-JUN-2002; 2002JP-00167239
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (SUMO ) SUMITOMO CHEM CO LTD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               17-OCT-2002; 2002WO-JP010789
                                                                                                     20-NOV-2002; 2002WO-US037411.
                                                                                                                       23-NOV-2001; 2001US-00021707.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
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                                                                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                         (ISIS-) ISIS PHARM INC
                                /*tag=
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Best Local Similarity
                                                                                                                                                            Dobie K;
                                                                                                                                                                           WPI; 2003-505193/47.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WO2003040370-A1
                                                                    WO2003046132-A2
                        modified base
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                                                                                                                                                                                                                                                                                                                                                           16;
                                                                                    05-JUN-2003
                                                                                                                                                            Karras JG,
                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADC36216
                                                                                                                                                                                                                                                                                                                                                                                                19
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                                                                                                                                                The invention relates to a novel DNA encoding a weed controller metabolism protein. A protein of the invention has herbicide activity. The proteins and their encoded genes are useful e.g. in constructing new breeds of herbicide-resistant plants and also in developing various agrochemicals. The present sequence is used in the exemplification of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Antisense; ss; human; CD81; TAPA-1; tetraspanin; viral infection; cocaine addiction; autoimmune disorder; antiinflammatory; antibacterial; virucide; antiparasitic; inflammatory disorder; parasitic infection; bacterial infection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /mod_base= OTHER
/note= "Phosphorothioate backbone and all cytidines are
-methyl cytidines"
Weed controller metabolism proteins deactivating porphyrinogen oxidase (PPO)-inhibiting herbicides by N-demethylation and their genes, useful e.g. in constructing new breeds of herbicide-resistant plants.
                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
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/mod_base= OTHER
/note= "2'-methoxyethyl nucleotide"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    note= "2'-methoxyethyl nucleotide'
                                                                                                                                                                                                                                                                                                                                Sequence 20 BP; 4 A; 5 C; 7 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human CD81/TAPA-1 antisense oligonucleotide #20.
                                                                                                                                                                                                                                                                                                                                                                            Score 14.2; DB 1;
Pred. No. 8.7e+02;
0; Mismatches 3;
                                                                                                            Disclosure; SEQ ID NO 83; 812pp; Japanese
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1222 GIGGAGGAACAGCIACACT 1240
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               base= OTHER
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 BP.
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                                                                                                                                                                                                                                                                                                                                                                                         0.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADC35560 standard; DNA; 20
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/*tag= b
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                                                                                                                                                                                                                                                                                                                                                                                                                                          16; Conservative
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/mod_bag
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The invention relates to a compound (antisense oligonucleotide) ... hybridising with the eighth nucleobase portion of an active site on a nucleic acid molecule encoding CD81 (Also known as TAPA-1, a tetraspanin) comprising the expression of CD81. Also included is a composition emposition the antisense oligonucleotide and a carrier or a diluent. The antisense oligonucleotide is useful for inhibiting the expression of CD81 in cells or tissues. The antisense oligonucleotide is also useful for treating infections preferably viral, bacterial and parasitic and diseases such as inflammatory disorders and autoimmune disorders. The
Novel compound hybridizing with nucleic acid molecule encoding CDB1 and inhibiting the expression of CDB1, useful for treating infections and disease associated with expression of CDB1 such as inflammation disorder.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /note= "Phosphorothioate backbone, All cytidine residues are 5-methylcytidines"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human; antisense; pituitary tumour-transforming gene; securin; cancer;
PTTG1; TUTR1; BSP-1 associated protein; gene therapy; phosphorothioate;
                                                                                                                                                                                                                                                                                                                                                       Gaps
                                                                                                                                                                                                                      disease or condition is characterised by chemical dependency (e.g. cocaine addiction). The present sequence is a CD81 antisense oligonucleotide of the invention.
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16. .20
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                                                                                                                                                                                                                                                                                                                  0.8%; Score 14.2; DB 1; Length 20; 34.2%; Pred. No. 8.7e+02;
                                                                                                                                                                                                                                                                                                                                                    3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human PTTG1 antisense oligonucleotide ISIS #131034.
                                                                                                                                                                                                                                                                                Sequence 20 BP; 4 A; 6 C; 4 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                0; Mismatches
                                                         Claim 3; SEQ ID NO 32; 55pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                           855 CAAGGACCTGAAGCAGTAC 873
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /mod_base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /mod_base= OTHER
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                                                                                                                                                                                                                                                                                                                                                                                                     19 CAAGGATGTGAAGCAGTTC 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAD62975 standard; DNA; 20 BP.
                                                                                                                                                                                                                                                                                                                                 84.2%;
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                                                                                                                                                                                                                                                                                                        Query Match
Best Local Similarity 84.25
Matches 16; Conservative
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/*tag= b
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modified_base
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                                                                                                                                                       invention relates to novel antisense compounds targetted to a nucleic
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                                                                                                                                                                     acid encoding (pituitary tumour-transforming gene) FTTGI (also known as securin, TUTRI and ESP-I associated protein) to inhibit its expression. Antisense compounds of the invention are useful for preparing compositions for treating pituitary, breast, colorectal, testicular, pulmonary and epithelial cancer. The invention is also used in gene therapy. The present sequence is an antisense oligonucleotide targetted to human PTTGI DNA. This sequence is used in the exemplification of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Novel polynucleotide useful for detecting single nucleotide polymorphisms
                                               New compound, having a sequence targeted to a nucleic acid encoding PTTG1, useful for preparing a composition for treating e.g. pitultary, colorectal, breast, testicular, pulmonary or epithelial cancers.
                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     human; single nucleotide polymorphism; microarray; side effect; ss;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Single nucleotide polymorphism detection primer, SEQ ID No 1779.
                                                                                                                                                                                                                                                                                                                                                                                                                .,
                                                                                                                                                                                                                                                                                                                                                                      0.8%; Score 14.2; DB 1; Length 20; 34.2%; Pred. No. 8.7e+02;
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                                                                                                                                                                                                                                                                                                                            Sequence 20 BP; 6 A; 6 C; 2 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                               84.2%; Preq. .v.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (KAGA-) KAGAKU GIJUTSU SHINKO JIGYODAN,
                                                                                                                                                                                                                                                                                                                                                                                                                                          1010 AGAGGGGAGAGCTCAAGCT 1028
                                                                                                                      Claim 3; Page 25; Opp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADF88196 standard; DNA; 20 BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
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Best Local Similarity 84.2
Matches 16; Conservative
                WPI; 2003-831618/77.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       primer; PCR.
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                                                                                                                                                                                                                                                                                               invention
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(first entry)

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Pharmaceutical composition for treating ailments associated with impaired respiration, has Oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or
                                                                                  Human; antisense; lung dysfunction; nasal airway dysfunction; antiinflammatory steroid; ubiquinone; antiinflammatory; antiallergic; antiasthmatic; hypotensive; immunosuppressive; cyfostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy; lung inflammation; respiratory disease; ds.
                                                                Human oligonucleotide sequence.
                                                                                                                                                                                                                                                                                                       24-APR-2001; 2001US-0286137P.
                                                                                                                                                                                                                                                                           23-APR-2002; 2002WO-US013135.
                                                                                                                                                                                                                                                                                                                                 (EPIG-) EPIGENESIS PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2003-229219/22
                                                                                                                                                                                                                        WO200285308-A2.
                                                                                                                                                                                              Homo sapiens.
                                       17-OCT-2003
                                                                                                                                                                                                                                                  31-OCT-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ubiquinone.
                                                                                                                                                                                                                                                                                                                                                                              Miller S,
              ABZ93135;
                                                                                                                                                                                                                                                                                                                                                                   Nyce JW,
 The present invention relates to a method for detecting microorganisms using primers (ADF90918-ADF91145). The method is used for detecting microorganisms (bacteria, fungi, protozoa, viruses) which cause diarrhoea symptoms, and pathogenic microbe (bod poisoning. The method can be used to detect unspecified microbes, or specific pathogens, or for the simultaneous detection of many kinds of microorganism.
                                                                                                                                           ..
diagnosis of disease and determination of side effect to a medical agent. The isolated human gene is also effective in detecting single nucleotide polymorphisms in a human gene. This polymucleotide sequence represents one of the PCR primers used in the single nucleotide polymorphism detection method of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Rapid, sensitive detection of specific or unspecified microbes causing diarrhea and food poisoning, using primers which target universal and specific genes, and amplifying by PCR under heat cycle conditions suitable for many detections.
                                                                                                                                             Gaps
                                                                                                                                                                                                                                                                                                                                                                                 Detection; microorganism; PCR; primer; bacterium; fungus; protozoan;
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14.2%; Pred. No. 8.7e+02;
ve 0; Mismatches 3; Indels
                                                                                                                0.8%; Score 14.2; DB 1; Length 20; 84.2%; Pred. No. 8.7e+02;
                                                                                                                                             Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 20 BP; 5 A; 6 C; 1 G; 8 T; 0 U; 0 Other;
                                                                                  Sequence 20 BP; 7 A; 7 C; 4 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                        Microorganism detection PCR primer, SEQ ID 72.
                                                                                                                                              0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                     virus; diarrhoea; food poisoning; ss
                                                                                                                                                                          1611 CTAAGCCACAGACCGAGGC 1629
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                                                                                                                                                                                                                                                                           ADF90989 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                  (first entry)
                                                                                                                     Query Match 0.8
Best Local Similarity 84.2
Matches 16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (RAKA-) RAKAN KK.
(GIFU-) GIFU DAIGAKUCHO.
                                                                                                                                                                                                                                                                                                                                                                                                                                   Listeria monocytogenes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2003-793230/75.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                             JP2003164282-A.
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                                                                                                                                                                                                                                                                                                        ADF90989;
                                                                                                                                                                                                                                                 RESULT 1067
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Katz E, Pabalan J, Aguilar D; S;

Li Y, Sandrasagra A, Tang L, Shahabuddin

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The invention relates to a novel pharmaceutical composition, which has a first active agent comprising an oligonucleotide antisense to the initiation codon, coding region, 5' or 3' end genomic flanking regions, comprising an genomic flanking regions, comparisons of second active agent comprising an associated within 2-10 nucleotides of junctions of genes encoding a polypeptide associated with lung and/or continifiammatory steroid and ubiquinone. A composition of the invention and as second active agent comprising an antisines gene therapy. The composition may have a immunosuppressive, and cytostatic activity. The composition may have a immunosuppressive, and cytostatic activity. The composition may have a composition are prophylactic or therapeutic respiratory effect of an for enhancing the prophylactic or therapeutic respiratory of an entiinflammatory steroid in a subject, for reducing or depleting levels of continional continuous surfactant in a subject, for reducing or depleting levels of inguinone or receptor, producing bronchodilation, increasing levels of ubiquinone or receptor, producing bronchodilation, increasing levels of ubiquinone or lung inflammation, lung allergies, or a respiratory disease or condition. Ung inflammation, but was obtained in electronic format directly from WIPO
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84.2%; Pred. No. 8.7e+02;
iive 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 20 BP; 8 A; 5 C; 3 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   at ftp.wipo.int/pub/published_pct_sequences
Disclosure; SEQ ID NO 8377; 872pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1521 GGAGATTCAGCTACAAAAG 1539
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GGAAATTCACCTTCAAAAG 19
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nes 16; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Matches
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ID ABZ8
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Gaps

; 0

1503 TTCCATATTTGCACTAAAG 1521

84.2%;

Query Match Best Local Similarity

16; Conservative

Matches

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TICCATCTTTCCACTAATG 20

BP.

ABZ93135 standard; DNA; 20

RESULT 1068

ABZ93135 ID ABZ9

Tue Nov

ABZ85058;

Human, antisense; lung dysfunction; nasal airway dysfunction, antiinflammatory steroid; ubiquinone; antiinflammatory; antiallergic; antiastimatic; hypotensive; immunosuppressive; cycostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy; lung inflammation; respiratory disease; ds. treating ailments associated with impaired Pharmaceutical composition for treating ailments associated with impaire respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or Katz E, Pabalan J, Aguilar D; Claim 15; SEQ ID NO 300; 872pp; English. Li Y, Sandrasagra A, Ka Tang L, Shahabuddin S; Human oligonucleotide sequence. 23-APR-2002; 2002WO-US013135. 24-APR-2001; 2001US-0286137P. (EPIG-) EPIGENESIS PHARM INC. 17-OCT-2003 (first entry) WPI; 2003-229219/22. WO200285308-A2. Homo sapiens. 31-OCT-2002 ubiquinone. Nyce JW, 1 Miller S,

The invention relates to a novel pharmaceutical composition, which has a first active agent comprising an oligonucleotide antisense to the confinitation codon, coding region, 5' or 3' end genomic flanking regions, 5' and 3' intron-exon junctions, or regions within 2-10 nucleotides of junctions of genes encoding a polypeptide associated with lung and/or nasal airway dysfunction and a second active agent comprising an continiflammatory steroid and ubiquinone. A composition of the invention has antiinflammatory, antiallergic, antiasthmatic, hypotensive, confirmunosuppressive, and cytostatic activity. The composition may have a cuse in antisense gene therapy. The composition is useful for treating or the preventing a respiratory, lung or mallignant disease or condition, also continiflammatory steroid in a subject; for reducing or depleting levels of of, or reducing the prophylactic or therapeutic respiratory effect of an antiinflammatory steroid in a subject; for reducing levels of adenosine of, or receptor, producing bronchodilation, increasing levels of ubiquinone or lung inflammation, lung allergies, or a respiratory disease or condition.

Condition in the sequence data for this patent is not represented in the printed secretic attention in the printed and the printed of the condition of the con ftp.wipo.int/pub/published_pct_sequences

Sequence 20 BP; 2 A; 4 C; 7 G; 7 T; 0 U; 0 Other;

Gaps .. 0.8%; Score 14.2; DB 1; Length 20; 34.2%; Pred. No. 8.7e+02; Ive 0; Mismatches 3; Indels 623 AGCTGGACAAACTGGGCGA 641 ilarity 84.2%; Conservative Query Match Best Local Similarity Matches 16; Conserv

ABZ85420 standard; DNA; 20 RESULT 1070 ABZ85420/ ID ABZ8

BP.

19 ACCTGAACAACTGGCCGA 1

qq à

Human; antisense; lung dysfunction, nasal airway dysfunction, antiinflammatory; sucroid; ubiquinone; antiinflammatory; antiallergic; antiasthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy; lung inflammation; respiratory disease; ds. Pabalan J, Sandrasagra A, Katz E, Human oligonucleotide sequence. 23-APR-2002; 2002WO-US013135. 24-APR-2001; 2001US-0286137P. (EPIG-) EPIGENESIS PHARM INC. (first entry) Li Y, Sar Tang L, WO200285308-A2. Homo sapiens. 17-0CT-2003 31-OCT-2002. ABZ85420; Miller S, Nyce JW,

Pharmaceutical composition for treating ailments associated with impaired

The invention relates to a novel pharmaceutical composition, which has a first active agent comprising an oligomuclectide antisense to the initiation codon, coding region, 5' or 3' end genomic flanking regions, 5' and 3' intron-exon junctions, or regions within 2-10 nucleotides of junctions of genes encoding a polypeptide associated with lung and/or nasal airway dysfunction and a second active agent comprising an antihillammatory steroid and ubiquinone. A composition of the invention antibiliammatory, antiallergic, antiasthmatic, hypotensive, immunosuppressive, and cytostatic activity. The composition may have a use in antisense gene therapy. The composition is useful for treating or preventing arespiratory, lung or malignant disease or condition, also for enhancing the prophylactic or therapeutic respiratory effect of an antiinflammatory steroid in a subject, for reducing levels of antiinflammatory steroid in a subject, for reducing levels of acceptor, producing bronchodilation, increasing levels of adenosine increasing levels of increasing i lung surfactant in a subject's tissue, or treating bronchoconstriction, lung inflammation, lung allergies, or a respiratory disease or condition. Note: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO at ftp.wipo.int/pub/published_pct_sequences

Sequence 20 BP; 4 A; 7 C; 8 G; 1 T; 0 U; 0 Other;

Gaps .. 0 Score 14.2; DB 1; Length 20; Pred. No. 8.7e+02; 3; Indels 0; Mismatches 84.2%; Local Similarity 84.2 les 16; Conservative Query Match Best Loca Matches

1029 GGCTGACTTTGGCCTGGCC 1047

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0

GGCTGCCCTTGACCTGGCC 1

19

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ABZ85267 standard; DNA; 20 BP ABZ85267/c Ω

Aguilar D;

WPI; 2003-229219/22.

respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or ubiquinone.

Claim 15; SEQ ID NO 662; 872pp; English.

RESULT 1071

Human; antisense; lung dysfunction; nasal airway dysfunction; antiallargic; antiallammatory steroid; ubiquinone; antiinflammatory; antiallargic; antiasthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy; inflammation; respiratory disease; ds Human oligonucleotide sequence. (first entry) 17-OCT-2003 ung

Homo sapiens.

WO200285308-A2.

31-OCT-2002

23-APR-2002; 2002WO-US013135.

24-APR-2001; 2001US-0286137P.

(EPIG-) EPIGENESIS PHARM INC

Aguilar D; Pabalan J, Katz E, ŝ Li Y, Sandrasagra A, Tang L, Shahabuddin Miller S, Ę, Nyce

WPI; 2003-229219/22.

Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or ubiquinone

Claim 15; SEQ ID NO 509; 872pp; English.

The invention relates to a novel pharmaceutical composition, which has a first active agent comprising an oligonucleotide antisense to the initiation codon, coding region, 5' or 3' end genomic flanking regions, continuity and a second active agent comprising an antistration sof genes encoding a polypeptide associated with lung and/or compariant and a second active agent comprising an antistral mamatory steroid and ubiquinone. A composition of the invention as antiinflammatory antiallergic, antiasthmatic, hypotensive, conse in antisense gene therapy. The composition may have a immunosuppressive, and cytostatic activity. The composition may have a immunosuppressive, and cytostatic activity. The composition may have a conse in antisense gene therapy. The composition may have a consent a respiratory, lung or malignant disease or condition, also continuially a respiratory. Lung or therapeutic respiratory effect of an for enhancing the prophylactic or therapeutic respiratory effect of an antisinal prophylactic or therapeutic respiratory disection or receptor, producing bronchodilation, increasing levels of adenosine or receptor, producing bronchodilation, increasing levels of ubiquinone or corporated time subject is tissue, or treating bronchoconstriction, congruent in a subject is tissue, or treating bronchoconstriction, but was obtained or a respiratory disease or condition. Specification, but was obtained in electronic format directly from WIPO at ftp.wipo.int/pub/published_pct_sequences

Sequence 20 BP; 4 A; 6 C; 6 G; 4 T; 0 U; 0 Other;

Gaps ·; 0.8%; Score 14.2; DB 1; Length 20; 34.2%; Pred. No. 8.7e+02; ive 0; Mismatches 3; Indels 84.2%; Query Match 0.8° Best Local Similarity 84.2 Matches 16; Conservative

8

ABZ84777 standard; DNA; 20 RESULT 1072 ABZ84777,

BP.

(first entry) 17-OCT-2003 ABZ84777;

Human oligonucleotide sequence.

Human; antisense; lung dysfunction; nasal airway dysfunction; antinflammatory steroid; ubiquinone; antiinflammatory; antiallergic; antiasthmatic; hypotensive; immunosuppressive; cyfostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy; lung inflammation; respiratory disease; ds.

Homo sapiens.

WO200285308-A2.

31-OCT-2002.

23-APR-2002; 2002WO-US013135.

24-APR-2001; 2001US-0286137P.

(EPIG-) EPIGENESIS PHARM INC.

Katz E, Pabalan J, Aguilar D; Sandrasagra A, Ka L, Shahabuddin S; Li Y, San Tang L, Nyce JW, I Miller S,

WPI; 2003-229219/22

Pharmaceutical composition for treating ailments associated with impaired respiration, has Oilgo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or ubiquinone

Claim 15; SEQ ID NO 19; 872pp; English.

The invention relates to a novel pharmaceutical composition, which has a first active agent comprising an oligonucleotide antisense to the initiation codon, coding region, 5' or 3' end genomic flanking regions, 5' and 3' intron-exon junctions, or regions within 2-10 nucleotides of junctions of genes encoding a polypeptide associated with lung and/or nasal airway dysfunction and a second active agent comprising an antiinflammatory steroid and ubiquinone. A composition of the invention and antisense gene therapy. The composition of the invention of preventing a respiratory, lung or malignant disease or condition, also for enhancing the prophylactic or therapeutic respiratory effect of an coff or reducing sensitivity to adenosine, reducing or depleting levels of or receptor, producing bronchodilation, increasing levels of adenosine or receptor, producing bronchodilation, increasing levels of ubiquinone or receptor, producing bronchodilation, increasing levels of ubiquinone or lung surfactant in a subject, stissue, or treating bronchoconstriction. Unug allergies, or a respiratory disease or condition. Successing the sequence date for this patent is not represented in the printed segment of the printed or presented in the printed or precent of the printed or presented in the printed or presented in the printed or precent or presented in the printed or presented in the printed or precent or presented in the printed or presented at ftp.wipo.int/pub/published_pct_sequences

Sequence 20 BP; 5 A; 3 C; 7 G; 5 T; 0 U; 0 Other;

Gaps .. 0 0.8%; Score 14.2; DB 1; Length 20; 84.2%; Pred. No. 8.7e+02; Live 0; Mismatches 3; Indels Best Local Similarity 84.2 Matches 16; Conservative Query Match

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993 GAACCTGCTCATCAACGAG 1011

ð g

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GAACCTGCTCATCTCCAAG 1 19

RESULT 1073 ABZ87947

ABZ87947 standard; DNA; 20 BP

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ABZ87947;
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17-OCT-2003 (first entry)

Human oligonucleotide sequence.

antiinflammatory steroid, ubiquinone, antiinflammatory, antiallergic, antiasthmatic, hypotensive, immunosuppressive, cytostatic, gene therapy, antisense gene therapy; respiratory; lung, adenosine sensitivity; adenosine receptor, bronchodilation; bronchodonstriction; lung allergy; lung inflammation; respiratory disease; ds. lung dysfunction; nasal airway dysfunction; antisense;

Homo sapiens.

WO200285308-A2.

31-OCT-2002.

23-APR-2002; 2002WO-US013135.

24-APR-2001; 2001US-0286137P.

(EPIG-) EPIGENESIS PHARM INC.

Pabalan J, Aguilar D; Katz E, Li Y, Sandrasagra A, Ka Tang L, Shahabuddin S; Miller S, Nyce JW,

WPI; 2003-229219/22.

Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or

Disclosure; SEQ ID NO 3189; 872pp; English.

The invention relates to a novel pharmaceutical composition, which has a first active agent comprising an oligomucleotide antisense to the first active agent comprising an oligomucleotide antisense to the initiation codon, coding region, 5' or 3' end genomic flanking regions, 5' and 3' intron-exon junctions, or regions within 2.0 nucleotides of nasal alraway dysfunctions, or regions within 2.0 nucleotides of nasal alraway dysfunction and a second active agent comprising an antiinflammatory steroid and ubiquinone. A composition of the invention has antiinflammatory, and cytostatic, antiachmatic, hypotensive, numbrous and cytostatic activity. The composition may have a serventing a respiratory, lung or malignant disease or condition, also preventing a respiratory, lung or malignant disease or condition, also antiinflammatory steroid in a subject, for reducing levels of adenosine of, or reducing sensitivity to adenosine, reducing levels of adenosine lung surfactant in a subject, stating levels of adenosine lung surfactant in a subject setsiang levels of ubiquinone or lung surfactant in a subject setsiang levels of ubiquinone or lung surfactant in a subject setsiang levels of upiquinone or lung surfactant in a subject setsiang levels of upiquinone or lung surfactant in a subject setsiang levels of upiquinone or lung surfactant in a subject setsiang levels of upiquinone or lung surfactant in a subject setsiang levels of upiquinone or lung surfactant in a subject setsiang levels of upiquinone or lung surfactant in a subject setsiang levels of upiquinone or lung surfactant in a subject setsiang levels of upiquinone or lung surfactant in a subject setsiang levels of upiquinone or lung surfactant in a subject setsiang levels of upiquinone or lung surfactant in a subject setsiang levels of upiquinone or lung surfactant in a subject setsiang levels of upiquinone or lung surfactant in a subject setsiang levels of upiquinone or lung surfactant in a subject setsiang levels of upiquinone or lung surfactant in a subject setsia lung inflammation, lung allergies, or a respiratory disease or condition. Note: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO ftp.wipo.int/pub/published_pct_sequences

Sequence 20 BP; 10 A; 2 C; 7 G; 1 T; 0 U; 0 Other;

Gaps ; 0.8%; Score 14.2; DB 1; Length 20; 34.2%; Pred. No. 8.7e+02; Ive 0; Mismatches 3; Indels 84.28; Conservative Local Similarity les 16; Conserv Query Match Best Loca Matches

1008 CGAGAGGGGAGGCTCAAG 1026

à g

CGAGAGAGAGATCAAG 19

ABZ87022 standard; DNA; 20 RESULT 1074 ABZ87022/c П

17-OCT-2003 (first entry) ABZ87022;

antiinflammatory steroid; uniquinone; antiinflammatory; antiallergic; antiasthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy; lung inflammation; respiratory disease; ds. lung dysfunction; nasal airway dysfunction; Human oligonucleotide sequence. Human; antisense;

Homo sapiens.

WO200285308-A2

31-OCT-2002,

23-APR-2002; 2002WO-US013135.

24-APR-2001; 2001US-0286137P.

(EPIG-) EPIGENESIS PHARM INC

Pabalan J, Aguilar D; Katz E, Sandrasagra A, Ka J, Shahabuddin S; Li Y, Sar Tang L, Nyce JW, Miller S,

WPI; 2003-229219/22.

Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or ubiquinone

Claim 15; SEQ ID NO 2264; 872pp; English.

first active agent comprising an oligomucleotide antisense to the initiation codon, coding region, 5' or 3' end genomic flanking regions, 5' and 3' intron-exon junctions, or regions within 2-10 nucleotides of unctions of genes encoding a polypeptide associated with lung and/or nasal airway dysfunction and a second active agent comprising and/or antinflammatory steroid and ubiquinone. A composition of the invention and antifflammatory, antiallergic, antiasthmatic, hypotensive, immunosuppressive, and cytostatic activity. The composition may have a use in antisense gene therapy. The composition is useful for treating or preventing a respiratory, lung or malignant disease or condition, also for enhancing the prophylactic or therapeutic respiratory effect of an antinflammatory steroid in a subject, for reducing or depleting levels of, or reducing sensitivity to adenosine, reducing levels of ubiquinone or increasing levels of upiquinone or increasing levels of ubiquinone or increasing levels of ubiquinone or increasing levels of upiquinone or increasing upicatione or increasing up lung inflammation, lung allergies, or a respiratory disease or condition. Note: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO The invention relates to a novel pharmaceutical composition, which has a lung surfactant in a subject's tissue, or treating bronchoconstriction ftp.wipo.int/pub/published_pct_sequences

Sequence 20 BP; 7 A; 4 C; 4 G; 5 T; 0 U; 0 Other;

·. Score 14.2; DB 1; Length 20; Pred. No. 8.7e+02; 0; Mismatches 3; Indels 0.8%; 16; Conservative Query Match Best Local Similarity Matches

1412

1394 CCAAGCTGTTGCAGTTTGA

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0

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Gaps

ccaagcrgargracrirga 1 13 RESULT 1075 g

ABZ88149 standard; DNA; 20 BP ABZ88149/ ID ABZ8

Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or antinflammatory steroid; ubiquinone; antinflammatory; antiallergic; antiathergic; antiathergic; antiathergic; pypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy; Aguilar D; antisense; lung dysfunction; nasal airway dysfunction; Katz E, Pabalan J, Sequence 20 BP; 3 A; 3 C; 8 G; 6 T; 0 U; 0 Other; at ftp.wipo.int/pub/published_pct_sequences lung inflammation; respiratory disease; ds. Disclosure; SEQ ID NO 3391; 872pp; English. Li Y, Sandrasagra A, Tang L, Shahabuddin Human oligonucleotide sequence. 23-APR-2002; 2002WO-US013135. 24-APR-2001; 2001US-0286137P. (EPIG-) EPIGENESIS PHARM INC (first entry) WPI; 2003-229219/22. WO200285308-A2. Homo sapiens. 17-0CT-2003 31-OCT-2002 Nyce JW, I Miller S, ubiquinone

The invention relates to a novel pharmaceutical composition, which has a first active agent comprising an oligonucleotide antisense to the initiation codon, coding region, 5' or 3' end genomic flanking regions, 5' or 3' end genomic flanking regions, 5' or 3' end genomic flanking regions, 5' and 3' intron-exon junctions, or regions within 2-10 nucleotides of junctions of genes encoding a polypeptide associated with lung and/or nasal airway dysfunction and a second active agent comprising an entiinflammatory steroid and ubiquinone. A composition of the invention has antiinflammatory, antiallergic, antiasthmatic, hypotensive, conse in antisense gene therapy. The composition may have a immunosuppressive, and cytostatic activity. The composition may have a conse in antisense gene therapy. The composition may have a preventing a respiratory, lung or mallignant disease or condition, also for enhancing the prophylactic or therapeutic respiratory effect of an antiinflammatory steroid in a subject, for reducing or depleting levels of, or reducing sensitivity to adenosine, reducing or depleting levels of, or receptor, producing bronchdilation, increasing levels of adenosine receptor, producing bronchdilation, increasing levels of ubiquinone or lung inflammation, lung allergies, or a respiratory disease or condition. Under the sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO

0.8%; Score 14.2; DB 1; Length 20; 34.2%; Pred. No. 8.7e+02; ve. 0; Mismatches 3; Indels 84.2%; 16; Conservative Query Match Best Local Similarity Matches 16; Conserv

8

ABZ87509 standard; DNA; 20 BP RESULT 1076 ABZ87509/c ID ABZ87

Human oligonucleotide sequence. (first entry) 17-0CT-2003

ABZ87509;

antiinflammatory steroid; ubiquinone; antiinflammatory; antiallergic; antiasthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy; antisense; lung dysfunction; nasal airway dysfunction; lung inflammation; respiratory disease; ds

Homo sapiens.

WO200285308-A2.

31-OCT-2002.

23-APR-2002; 2002WO-US013135.

24-APR-2001; 2001US-0286137P.

(EPIG-) EPIGENESIS PHARM INC.

Katz E, Pabalan J, Aguilar D; Sandrasagra A, Ka L, Shahabuddin S; Li Y, San Tang L, Miller S, Nyce JW,

WPI; 2003-229219/22.

Pharmaceutical composition for treating ailments associated with impaired respiration, has Oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or ubiquinone

Disclosure; SEQ ID NO 2751; 872pp; English.

The invention relates to a novel pharmaceutical composition, which has a first active agent comprising an oligonucleotide antisense to the initiation codon, coding region, 5' or 3' end genomic flanking regions, or regions within 2-10 nucleotides of 5' and 3' intron-exon junctions, or regions within 2-10 nucleotides of junctions of genes encoding a polypeptide associated with lung and/or antihifammatory steroid and ubiquinone. A composition of the invention has antihifammatory; antiallargic, antiasthmatic, hypotensive, and cytostatic activity. The composition may have a communosuppressive, and cytostatic activity. The composition may have a conservation of preventing a respiratory, lung or malignant disease or condition, also preventing the prophylactic or therapeutic respiratory effect of an antihifammatory steroid in a subject, for reducing levels of adenosine of or receptor, producing bronchodilation, increasing levels of adenosine or lung surfactant in a subject, stissue, or treating bronchoconstriction, lung a surfactant in a subject stassue, or treating bronchoconstriction, lung a surfactant in a subject stassue, or treating bronchoconstriction, lung a surfactant in a subject, stissue, or treating bronchoconstriction, lung a surfactant in a subject, stissue, or treating bronchoconstriction, is not represented in the printed lung inflammation, lung allergies, or a respiratory disease or condition. Note: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO at ftp.wipo.int/pub/published_pct_sequences

Sequence 20 BP; 2 A; 6 C; 2 G; 10 T; 0 U; 0 Other;

Gaps ö 0.8%; Score 14.2; DB 1; Length 20; 44.2%; Pred. No. 8.7e+02; ve 0; Mismatches 3; Indels 84.2%; Best Local Similarity 84.2 Matches 16; Conservative Query Match

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733 715 CTGGAACATGAAGAGGGGG CTGGAACATGAAGAAGAG 20

> g à

. 0

Gaps

; 0

ABV77015 standard; DNA; 20 RESULT 1077 ABV77015,

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New internal transcribed spacer-derived oligonucleotide primer useful for detecting fungal pathogens such as Colletotrichum acutatum, Alternaria spp. or Cladosporium carpophilum.
                                                                                                                                                                                                                                                                                                                                                                                                                                              particular (175) region. The primers are useful for detecting a fungal pathogen such as Colletotrichum acutatum, Alternaria spp. or Cladosporium carpophilum. The primers are useful for detecting specific isolates of fungal pathogens and for monitoring disease development in plant populations, for assessing potential damage in a specific crop variety/pathogen strain relationship, for providing detailed information on the development and spread of specific pathogen races over extended geographical areas, and for detecting diseases with long latent phase
                                                                                                                                                                                                                                                                                                                                                                                                         PCR primers ABV77013-16 represent conserved primers designed for amplification of the fungal nuclear ribosomal RNA internal transcribed
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Guignardia, pathogen, internal transcribed spacer; ITS; citrus fruit intergenic sequence, intronic sequence, calmodulin, chitin synthase; citrus blackspot; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Guignardia internal transcribed spacer (ITS) reverse primer #2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Differentiating pathogenic and non-pathogenic Guignardia sp.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0.8%; Score 14.2; DB 1; Length 20;
84.2%; Pred. No. 8.7e+02;
tive 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 20 BP; 2 A; 6 C; 5 G; 7 T; 0 U; 0 Other;
                                                                                                                                                          SYGN ) SYNGENTA PARTICIPATIONS AG.
                                                                                                                                                                                                                                                                                                                                                                     Example 6; Page 20; 51pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1549 CTTCGGTCTTCGTCGATGC 1567
                                                                                                                                                                                                         Perry CV;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               BP
                                                       08-MAR-2002; 2002WO-EP002581
                                                                                                09-MAR-2001; 2001US-0274540P
                                                                                                                       24-AUG-2001; 2001US-00939379
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         09-OCT-2001; 2001US-0327982P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ACA61050 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                                                                                       Barnett CJ,
                                                                                                                                                                                                                                           WPI; 2003-092859/08.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2003-372133/35.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Best Local Similarity
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                 03-OCT-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            14-JUL-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           16;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Guignardia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Carroll GC;
                                                                                                                                                                                                    Beck JJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ACA61050;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PCR primers ABV77013-16 represent conserved primers designed for amplification of the fungal nuclear ribosomal RNA internal transcribed spacer (ITS) region. The primers are useful for detecting a fungal pathogen such as Colletotrichum acutatum, Alternaria spp. or Cladosporium carpophilum. The primers are useful for detecting specific isolates of fungal pathogens and for monitoring disease development in plant populations, for assessing potential damage in a specific crop variety/pathogen strain relationship, for providing detailed information on the development and spread of specific pathogen races over extended geographical areas, and for detecting diseases with long latent phase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New internal transcribed spacer-derived oligonucleotide primer useful for detecting fungal pathogens such as Colletotrichum acutatum, Alternaria spp. or Cladosporium carpophilum.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
                                                                                                                          Internal transcribed spacer region; ITS region; fungal pathogen;
Colletotrichum acutatum; Alternaria; Cladosporium carpophilum; PCR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Internal transcribed spacer region; ITS region; fungal pathogen;
Colletotrichum acutatum; Alternaria; Cladosporium carpophilum; PCR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ..
0
                                                                                   Primer ITS3 used to amplify fungal nuclear rDNA ITS region.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0.8%; Score 14.2; DB 1; Length 20;
84.2%; Pred. No. 8.7e+02;
Live 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Primer ITS2 used to amplify fungal nuclear rDNA ITS region.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 20 BP; 7 A; 5 C; 6 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                     (SYGN ) SYNGENTA PARTICIPATIONS AG.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example 6; Page 20; 51pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1549 CTTCGGTCTTCGTCGATGC 1567
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                                                                                                                                                                                                                                                                                                                                   08-MAR-2002; 2002WO-EP002581.
                                                                                                                                                                                                                                                                                                                                                                         09-MAR-2001; 2001US-0274540P.
24-AUG-2001; 2001US-00939379.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Beck JJ, Barnett CJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2003-092859/08.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Local Similarity
                                                                                                                                                                                                                                                    WO200277293-A2.
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                                            03-MAR-2003
                                                                                                                                                                                                                                                                                            03-OCT-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   16;
                                                                                                                                                                      primer; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      03-MAR-2003
                                                                                                                                                                                                           Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              primer; ss
    ABV77015;
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WO2003031933-A2
                                                                  Guignardia sp.
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 probes based
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S
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                         Guignardia
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                                                   ACA61051
                                                                                     Carroll
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Matches
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                                                                                                                                                                                                               The invention decsribes a method of differentiating pathogenic and non-pathogenic species of Guignardia (I). The method comprises obtaining a DNA sample from a citrus fruit infected with (I), immobilising the DNA, probing the immobilised DNA with a probe based on intergenic sequences and intronic sequences from within the calmodulin and chitin synthase genes, and demonstrating hybridisation with the probes to represent the pathogenic species and non-pathogenic species. The method is specific, rapid and useful for differentiating pathogenic species (e.g. diugnardia citricarpa, the causative agent of citrus blackspot) from non-pathogenic species of Guignardia. This sequence represents a primer used to isolate an internal transcribed spacer to allow characterisation of pathogenic
hybridization between DNA from Guignardia- infected citrus and sed on intronic sequences from calmodulin and chitin synthase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
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Pred. No. 8.7e+02;
0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 20 BP; 2 A; 6 C; 5 G; 7 T; 0 U; 0 Other;
                                                                                                                                                           Example 1; Page 19; 37pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0.8%;
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Best Local Similarity
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Guignardia internal transcribed spacer (ITS) forward primer #3.
ACA61051 standard; DNA; 20
                                                                                                                                  14-JUL-2003 (first entry)
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Guignardia: parhogen; internal transcribed spacer; ITS; citrus fruit; intergenic sequence; intronic sequence; calmodulin; chitin synthase; citrus blackspot; PCR; primer; ss.

09-OCT-2002; 2002WO-US032227

09-OCT-2001; 2001US-0327982P

(UYOR-) UNIV OREGON

WPI; 2003-372133/35.

Differentiating pathogenic and non-pathogenic Guignardia sp., by assessing hybridization between DNA from Guignardia- infected citrus and probes based on intronic sequences from calmodulin and chitin synthase genes.

Example 1; Page 20; 37pp; English.

The invention decsribes a method of differentiating pathogenic and non-pathogenic species of Guignardia (I). The method comprises obtaining a DNA sample from a citrus fruit infected with (I), immobilising the DNA, probing the immobilised DNA with a probe based on intergenic sequences and intronic sequences from within the calmodulin and chitin synthase genes, and demonstrating hybridisation with the probes to represent the

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New alpha- conotoxin-like peptides that inhibit the activity of neuronal nicotinic acetylcholine receptor, useful for treating stroke, pain, schizophrenia, Parkinson's disease, small cell lung carcinoma or
              rapid and useful for differentiating pathogenic species (e.g. Guignardia citricarpa, the causative agent of citrus blackspot) from non-pathogenic species of Guignardia. This sequence represents a primer used to isolate an internal transcribed spacer to allow characterisation of pathogenic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      neuroleptic; antiparkinsonlan; cytostatic; nootropic; neuroprotective; neuronal nicotinic acetylcholine receptor; nAChR; inhibitor; stroke; pain; cancer related pain; post-surgical pain; oral pain; estreminal neuralgia; post-surgical pain; oral pain; phantom limb pain; fibromyalgia; referrentic neuralgia; retematoid arthritis; inflammatory arthritis; neurogenic pain; reumatoid arthritis; inflammatory arthritis; neurogenic pain; parkinson's disease; small cell lung carcinoma; Alzheimer's disease; nerve injury; PCR; primer; ss.
pathogenic species and non-pathogenic species. The method is specific,
                                                                                                                                                                                       Gaps
                                                                                                                                                                                       ·,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Alpha-conotoxin; cerebroprotective; analgesic; anticonvulsant;
                                                                                                                                                    Length 20;
                                                                                                                                                                                     3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                          PCR primer for the isolation of peptide Vcl.1 #SEQ ID 5.
                                                                                                                    BP; 7 A; 5 C; 6 G; 2 T; 0 U; 0 Other;
                                                                                                                                                     Score 14.2; DB 1;
Pred. No. 8.7e+02;
                                                                                                                                                                                         0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 18; Page 31; 87pp; English.
                                                                                                                                                                                                                          1549 CTTCGGTCTTCGTCGATGC 1567
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gayler K,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             29-MAR-2001; 2001AU-00004094.
                                                                                                                                                     0.8%;
                                                                                                                                                                                                                                                            CIGCGIICTICATCGAIGC
                                                                                                                                                                                                                                                                                                                                               ABZ21316 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
                                                                                                                                                                                          Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Alzheimer's disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2003-103260/09
                                                                                                                                                                           Similarity
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KHALIL Z.
GAYLER K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conus victoriae.
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                                                                                                                         Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                  24-FEB-2003
                                                                                                                                                                                          16;
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                                                                                           Guignardia
                                                                                                                                                                                                                                                                                                                                                                                  ABZ21316;
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(KHAL/)
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                                                                                                                                                                                                                                                             13
                                                                                                                                                       Query Match
                                                                                                                                                                              Local
                                                                                                                                                                                                                                                                                                               RESULT 1081
                                                                                                                                                                                          Matches
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dental pain, referred trigeminal neuralgia, post-herpetic neuralgia, phantom limb pain, fibromyalgia, reflex sympathetic dystrophy, pain associated with inflammatory conditions, rheumatoid arthritis or inflammatory arthritis, or pain resulting from conditions associated with neurogenic or neuropathic pain), epilepsy, nicotine addiction, schizophrenia, Parkinson's disease, small cell lung carcinoma, or Alzheimer's disease. The alpha-conotoxin-like peptide is also useful for accelerating recovery from nerve injury. The peptides are also useful as research reagents for investigating nicotinic acetylcholine receptor physiology and pharmacology. The current sequence represents a PCR primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The present invention relates to a method of finding at least one murine heavy chain variable region (VH)-encoding polynucleotide encoding at least the VH of an antibody having binding specificity for tumourassociated sialylated glycoprotein antigen (TAG-72). The method involves providing at least one test medium containing at least one murine anti-TAG-72 vH-encoding polynucleotide comprising at least one 5'-non-translated region (5'-NYR) of a murine anti-TAG-72 vH-encoding gene. The process is useful in finding at least one murine vH-encoding gene. The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Heavy chain variable region; VH;
tumour-associated sialylated glycoprotein antigen; TAG-72; therapy;
cancer; diagnosis; gene therapy; cytostatic; mouse; VH alpha TAG DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Finding polynucleotide encoding an antibody against tumor-associated glycoprotein antigen-72 (TAG-72), for treating cancer, by providing a test medium with murine anti-TAG-72 heavy variable region-encoding polynucleotide.
                                                                                                                                                                                                                                                               0.8%; Score 14.2; DB 1; Length 20;
84.2%; Pred. No. 8.7e+02;
ive 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mouse VH alpha TAG DNA specific primer, B72.3/CC92 HC.
                                                                                                                                                                                                                        Sequence 20 BP; 4 A; 2 C; 8 G; 6 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example; SEQ ID NO 44; 120pp; English.
                                                                                                                                                                 physiology and pharmacology. The cr
for the isolation of peptide Vc1.1
                                                                                                                                                                                                                                                                                                                                        889 AACATCATCAACATGCACA 907
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88US-00261942.
89US-00424362.
93US-00040687.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
                                                                                                                                                                                                                                                                                                   Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (DOWC ) DOW CHEM CO.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2003-851359/79.
                                                                                                                                                                                                                                                                                Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         19-OCT-1988;
24-OCT-1988;
19-OCT-1989;
31-MAR-1993;
24-MAR-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mus musculus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 US6641999-B1
                                                                                                                                                                                                                                                                                                 16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            06-MAY-2004
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADK66856;
                                                                                                                                                                                                                                                                 Query Match
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The present invention relates to a method of finding at least one murine heavy chain variable region (VH)-encoding polynucleotide encoding at least the VH of an antibody having binding specificity for tumourlessociated sialylated glycoprotein antigen (TAG-72). The method involves providing at least one test medium containing at least one murine anti-tag-72 VH-encoding polynucleotide comprising at least one 5'-nontragion (5'-NTR) of a murine anti-TAG-72 VH-encoding gene. The process is useful in finding at least one murine VH-encoding gene. The process is useful in finding at least the VH of an antibody having binding specificity for TAG-72. The antibodies against TAG-72 and compositions are useful in treating cancer, in in vivo diagnostic assays, in vivo therapy or radioinmunoguided surgery. The invention is also useful in gene therapy. The present sequence is mouse B72.3/CC92 immunoglobulin heavy chain DNA specific primer used in the exemplification of the
polynucleotide encoding at least the VH of an antibody having binding specificity for TAG-72. The antibodies against TAG-72 and compositions are useful in treating cancer, in in vivo diagnostic assays, in vivo therapy or radicimmunoguided surgery. The invention is also useful in gene therapy. The present sequence is mouse VH alpha TAG DNA specific primer used in the exemplification of the invention.
                                                                                                                                                                                                               Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Finding polynucleotide encoding an antibody against tumor-associated glycoprotein antigen-72 (TAG-72), for treating cancer, by providing test medium with murine anti-TAG-72 heavy variable region-encoding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Heavy chain variable region; VH;
tumour-associated sialylated glycoprotein antigen; TAG-72; therapy;
cancer; diagnosis; gene therapy; cytostatic; mouse; primer; ss.
                                                                                                                                                                                                               ..
0
                                                                                                                                                                         Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mouse B72.3/CC92 antibody heavy chain DNA specific primer,
                                                                                                                                                                   0.8%; Score 14.2; DB 1; Length 2
34.2%; Pred. No. 8.7e+02;
Ive 0; Mismatches 3; Indels
                                                                                                                                Sequence 20 BP; 4 A; 6 C; 2 G; 8 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Anderson WHK;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example; SEQ ID NO 24; 120pp; English.
                                                                                                                                                                                                                                            1293 GTCCAACGAGGAGTTCAAG 1311
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Rixon MW,
                                                                                                                                                                                                                                                                                ~
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                88US-00259943.
88US-00261942.
89US-00424362.
93US-00040687.
                                                                                                                                                                                                                                                                                                                                                                             BP.
                                                                                                                                                                                                                                                                              20 GTACAATGAGAAGTTCAAG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           14-FEB-2000; 2000US-00503653.
                                                                                                                                                                                      84.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         97US-00823105
                                                                                                                                                                                                                                                                                                                                                                         ADK66836 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                (first entry)
                                                                                                                                                                                                           Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gourlie B,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (DOWC ) DOW CHEM CO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2003-851359/79.
                                                                                                                                                                                      Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               polynucleotide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              19-0CT-1988;
24-0CT-1988;
19-0CT-1989;
31-MAR-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mus musculus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  US6641999-B1.
                                                                                                                                                                                                         16;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         24-MAR-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         34-NOV-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mezes PS,
                                                                                                                                                                     Query Match
                                                                                                                                                                                                                                                                                                                                                                                                             ADK66836;
                                                                                                                                                                                                                                                                                                                                        RESULT 1083
                                                                                                                                                                                        Best Loca
Matches
                                                                                                                                                                                                                                                                                                                                                       ADK66836/
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reduce the amount of target polypeptide present in the lungs. The

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                                                                                                                                                                                                                                                                                                                                                       respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antiinflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory disease; pulmonary vasoconstriction; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis; pulmonary transplantation rejection; ss; primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                This invention describes a novel composition (a) a first active agent,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Ç
                                                                                    Gaps
                                                                                                                                                                                                                                                                                                                                          antisense; bronchoconstriction; allergy; hyposecretion; pain;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Pharmaceutical composition for treating asthma, has antisense oligonucleotide containing less percentage of adenosine, targeted t nucleic acids associated with lung airway or lung dysfunction, and bronchodilating agent.
                                                                                   ..
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 E, Pabalan J, Aguilar
                                                        Length 20;
                                                                                   Indels
                         BP; 4 A; 6 C; 2 G; 8 T; 0 U; 0 Other;
                                                   Score 14.2; DB 1;
Pred. No. 8.7e+02;
0; Mismatches 3;
                                                                                                                                                                                                                                                                                                                A1672565-derived oligonucleotide SEQ ID 3391.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 15; SEQ ID NO 3391; 763pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Katz
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 s,
                                                                                                                  GICCAACGAGGAGTICAAG 1311
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sandrasagra A,
                                                                                   ;
0
                                                                                                                                              GIACAATGAGAAGITCAAG 2
                                                                                                                                                                                                                         BP
                                                        0.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           23-APR-2002; 2002WO-US013143.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       24-APR-2001; 2001US-0286036P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (EPIG-) EPIGENESIS PHARM INC
                                                                                                                                                                                                                         ABD24379 standard; DNA; 20
                                                                                                                                                                                                                                                                                   (first entry)
                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Li Y, Sar
Tang L,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2003-093058/08.
                                                                   Local Similarity
nes 16; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WO200285309-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Homo sapiens
                                                                                                                                                                                                                                                                                     29-JUL-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             31-OCT-2002.
                            Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Nyce JW, I
Miller S,
invention
                                                                                                                                                                                                                                                       ABD24379;
                                                                                                                    1293
                                                                                                                                                 20
                                                          Query Match
                                                                                                                                                                                             RESULT 1084
                                                                                       Matches
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The invention also describes a kit, that comprises: (a) a delivery device, in separate containers, (b) the oligomocleotides, (c) instructions for adding a carrier and for use of the kit. The composition of the invention has antiallersic, antiinflammatory, antiasthmatic, analgesic, hypotensive, immunosuppressive and cytostatic activity, is a beta-adrenergic agonist. The composition is useful for preventing or treating a respiratory, lung or malignant disease. The administered composition comprises oligo and is administered to reduce the production or availability, or to increase the degradation of the target mRNA or to

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   pulmonary obstruction, and/or bronchoconstriction and/or lung pulmonary obstruction, and/or bronchoconstriction and/or lung inflammation, allergies and/or bronchoconstriction and/or lung inflammation, allergies, asthma, impeded respiration, respiratory unflammation, allergies, asthma, impeded respiration, respiratory distress syndrome, pain, cystic fibrosis, allergic rhinitis, pulmonary hypertension, emphysema, chronic obstructive pulmonary disease, pulmonary transplantation rejection, pulmonary infections, bronchis or cancer. The reduced adenosine content of the anti-sense oligos corresponding to thymidines present in the target RNA serves to prevent the breakdown of the oligonucleotides into products that free adenosine into the system e.g., lung, brain, heart, kidney, etc, tissue environment and thereby, to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             comprising oligonucleotides, effective for alleviating bronchoconstriction, respiratory tract inflammation, allergies and reducing adenosine sensitivity, levels of adenosine (A) or (A) receptors, surfactant depletion or hyposecretion, when administered to a mammal. The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antiinflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cycotatic; cystic fibrosis; respiratory disease; pulmonary vasoconstriction; respiratory disease; pulmonary vasoconstriction; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis; pulmonary transplantation rejection; ss; primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            This invention describes a novel composition (a) a first active agent
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Ç
                                                                                                                                                                                                                                                                                                                                Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Pharmaceutical composition for treating asthma, has antisense oligonucleotide containing less percentage of adenosine, targeted trucleic acids associated with lung airway or lung dysfunction, and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Katz E, Pabalan J, Aguilar D;
                                                                                                                                                                                                                                                                                                                                ..
                                                                                                                                                                                                                                                                                         Length 20;
                                                                                                                                                                                                                                                                               0.8%; Score 14.2; DB 1; Lengtu 2.84.2%; Pred. No. 8.76+02; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               S100 calcium binding protein A2-derived oligo SEQ ID 662
                                                                                                                                                                                                                                                        Sequence 20 BP; 3 A; 3 C; 8 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 15; SEQ ID NO 662; 763pp; English.
                                                                                                                                                                                                                   prevent any unwanted effects due to it
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ŝ
                                                                                                                                                                                                                                                                                                                                                                     994 AACCTGCTCATCAACGAGA 1012
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sandrasagra A,
5, Shahabuddin
                                                                                                                                                                                                                                                                                                                                                                                                         ACCCTGCTCATCAGCAAGA 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          23-APR-2002; 2002WO-US013143.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             24-APR-2001; 2001US-0286036P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ABD21650 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
                                                                                                                                                                                                                                                                                                                                     Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Li Y, Sar
Tang L,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2003-093058/08.
                                                                                                                                                                                                                                                                                                            Local Similarity
nes 16; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WO200285309-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                29-JUL-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     31-OCT-2002
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← S,
                                                                                                                                                                                                                                                                                                                                                                                                             13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ABD21650;
                                                                                                                                                                                                                                                                                                 Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Miller
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RESULT 1085
                                                                                                                                                                                                                                                                                                                                  Matches
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530

or lung

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The invention also describes a kit, that comprises: (a) a delivery device, in separate containers, (b) the oligomucleotides, (c) instructions for adding a carrier and for use of the kit. The composition of the invention has antiallergic, antiinflammatory, antiasthmatic, analgesic, hypotensive, immunosuppressive and cytostatic activity, is a ceta-adrenation equivalent of composition is useful for preventing or treating a respiratory, lung or malignant disease. The administered composition comprises oligo and is administered to reduce the production or availability, or to increase the degradation of the target mRNA or to reduce the amount of target polypeptide present in the lungs. The pulmonary obstruction, and/or bronchoconstriction and/or lung with a disease or condition such as pulmonary vasoconstriction, with a disease or condition such as pulmonary vasoconstriction, allergies, asthma, impeded respiration, respiratory when the complex condition such as pulmonary vasoconstriction, and condition such as pulmonary vasoconstriction, and condition such as pulmonary vasoconstriction, when the condition condition such as pulmonary vasoconstriction, where the amount of target fibrosis, allergies in cystic fibrosis, allergies, pulmonary when the condition are such as pulmonary experience.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         the oligonucleotides into products that free adenosine into the system e.g., lung, brain, heart, kidney, etc, tissue environment and thereby, to prevent any unwanted effects due to it
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     hypertension, emphysema, chronic obstructive pulmonary disease, pulmonary
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       transplantation rejection, pulmonary infections, bronchitis or cancer. The reduced adenosine content of the anti-sense oligos corresponding to thymidines present in the target RNA serves to prevent the breakdown of
                          expression of a target polypeptide associated with lung airway or lung dystunction or cancer and can be anti-sense to the corresponding mRNA.
oligonucleotides are derived from a gene encoding or regulating
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Sequence 20 BP; 4 A; 7 C; 8 G; 1 T; 0 U; 0 Other;

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Gaps
                         0;
0.8%; Score 14.2; DB 1; Length 20; 84.2%; Pred. No. 8.7e+02; ive 0; Mismatches 3; Indels
                                          1029 GGCTGACTTTGGCCTGGCC 1047
                    0;
                                                               Н
                                                               GGCTGCCTTGACCTGGCC
                        Conservative
          Similarity
   Ma.
Local S...
16;
                                                               19
 Query Match
                     Matches
                                          ò
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AA001432-derived oligonucleotide SEQ ID 8377. ABD29365 standard; DNA; 20 BP (first entry) 29-JUL-2004 ABD29365; RESULT 1086 4BD29365

analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory distress syndrome; allergic rhinitis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis; Human, antisense, bronchoconstriction, allergy, hyposecretion, pain, respiratory tract inflammation, adenosine sensitivity, lung, cancer, surfactant depletion, antiallergic, antiinflammatory, antiasthmatic, emphysema; chronic obstructive pulmonary disease pulmonary transplantation rejection; ss; primer.

Homo sapiens.

WO200285309-A2.

31-OCT-2002

23-APR-2002; 2002WO-US013143.

24-APR-2001; 2001US-0286036P.

(EPIG-) EPIGENESIS PHARM INC.

Aguilar D; Pabalan J, Katz E, Sandrasagra A, Ka L, Shahabuddin S; Li Y, Sar Tang L, Nyce JW, I Miller S,

WPI; 2003-093058/08.

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This invention describes a novel composition (a) a first active agent, comparising oligomuclectides, effective for alleviating bromchoconstriction, respiratory tract inflammation, allergies and bromchoconstriction, respiratory tract inflammation, allergies and reductional adenosine early propered to adenosine (A) or (A) receptors, surfactant depletion or hyposecretion, when administered to a mammal. The oligomuclectides are derived from a gene encoding or regulating cyperide associated with lung airway or lung dysfunction or cancer and can be anti-sense to the corresponding mRNA. The invention also describes a kit, that comprises: (a) a delivery or lung dysfunction or cancer and can be anti-sense to the corresponding mRNA. The invention has antiallargic, antiinflammacory, antiasthmatic, instructions for adding a carrier and for use of the kit. The composition of the invention has antiallargic, antiinflammacory, antiasthmatic, of the invention has antiallargic, antiinflammacory, antiasthmatic, beta-adrenergic agonist. The composition is useful for preventing or composition comprises oligo and is administered to reduce the production or availability, or to increase the degradation of the target mRNA or to reduce the amount of target polypeptide present in the lungs. The culmonary obstruction, and/or bronchoconstriction and/or lung inflammation, allergies and/or surfactant hypoproduction are associated with a disease or condition such as pulmonary vascocnstriction, with a disease or condition such as pulmonary vascocnstriction, with a disease or condition such as pulmonary vascocnstriction, thypertension, emphysema, chronic obstructive pulmonary disease, pulmonary infections, bronchitis or cancer. The reduced adenosine content of the anti-sense oligops corresponding to thymidines present in the target RNA serves to prevent the breakdown of the oligonuclectides into products that free adenosine into the system of egg., the oligonuclectides into the interest in the interest and the interest and the interest and the 
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                                                                                                                                                                                                                    describes a novel composition (a) a first active agent,
                                           5
Pharmaceutical composition for treating asthma, has antisense oligonuclectide containing less percentage of adenosine, targeted trucleic acids associated with lung airway or lung dysfunction, and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 14.2; DB 1; Length 20; Pred. No. 8.7e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 20 BP; 8 A; 5 C; 3 G; 4 T; 0 U; 0 Other;
                                                                                                                                                      SEQ ID NO 8377; 763pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    prevent any unwanted effects due to it
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 84.2%;
                                                                                             bronchodilating agent.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Local Similarity
                                                                                                                                                                                                                    invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
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Gaps 0 3; Indels 0; Mismatches 16; Conservative Matches

1521 GGAGATTCAGCTACAAAG 1539 GGAAATTCACCTTCAAAAG 19 ð g

29-JUL-2004 (first entry) ABD23252;

ABD23252 standard; DNA; 20 BP.

RESULT 1087

Human myosin X-derived oligonucleotide SEQ ID 2264.

Human; antisense; bronchoconstriction; allergy; hyposecretion; pain; respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antialinery, inflammatic; antialinflammaticy; antiasthmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory distress syndrome; allergic fibritis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis; pulmonary transplantation rejection; ss; primer.

Homo sapiens,

WO200285309-A2

31-OCT-2002.

10017621-381.rng

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This invention describes a novel composition (a) a first active agent, comprising oligonuclectides, effective for alleviating comprising oligonuclectides, effective for alleviating comprising oligonuclectides, respiratory tract inflammation, allergies and reducing adenosine sensitivity, levels of adenosine (A) or (A) receptors, surfactant depletion or hyposecretion, when administered to a mammal. The oligonuclectides are derived from a gene encoding or regularing dysfunction or cancer and can be anti-sense to the corresponding mRNA. The invention also describes a kit, that comprises: (a) a delivery device, in separate containers, (b) the oligonuclectides, (c) instructions for adding a carrier and for use of the kit. The composition of the invention has antiallergic, antiinflammatory, antiakfammatory, antiakfammatory, antiakfammatory, antiakfammatory, antiakfammatory, or onergonist. The composition is useful for preventing or treating a respiratory, lung or malignant disease. The administered composition comprises oligo and is administered to reduce the production or availability, or to increase the degradation of the target mRNA or to reduce the amount of target polypeptide present in the lungs. The cumposition and/or bronchoconstriction and/or lung inflammation, allergies and/or bronchoconstriction and/or lung inflammation, allergies and/or bronchoconstriction and/or inflammation, allergies and/or bronchoconstriction and/or inflammation, allergies and/or bronchoconstriction and/or treating a respiratory distress syndrome, pain, cystic fibrosis, allergic rhinitis or cancer. The reduced adenosine content of the anti-sense oligons corresponding to the rangel RNA serves to prevent the breakdown of the oligonuclectication beatt, kidney, etc, tissue environment and thereby, to try bronchocouse that the environment and thereby, to the product of the condition of the toit the delicest adenosine content of the antiance of the oligonuclectides into product that the delices of the oligonuclection and the toit transplanta
                                                                                                                                                                                                                                                                                                           Pharmaceutical composition for treating asthma, has antisense oligonucleotide containing less percentage of adenosine, targeted to nucleic acids associated with lung airway or lung dysfunction, and
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 15; SEQ ID NO 2264; 763pp; English.
                      23-APR-2002; 2002WO-US013143.
                                                                             24-APR-2001; 2001US-0286036P
                                                                                                                               (EPIG-) EPIGENESIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                      bronchodilating agent.
                                                                                                                                                                                                                                                                 WPI; 2003-093058/08.
                                                                                                                                                                                                          Miller S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Matches
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ó Gaps . 0 Score 14.2; DB 1; Length 20; Pred. No. 8.7e+02; 3; Indels Sequence 20 BP; 7 A; 4 C; 4 G; 5 T; 0 U; 0 Other; 0; Mismatches 0.8%; 84.2%; 16; Conservative Query Match Best Local Similarity à

a

ABD24177 standard; DNA; 20 BP 29-JUL-2004 ABD24177; RESULT 1088

(first entry)

Human calmodulin 2-derived oligonucleotide SEQ ID 3189.

Human; antisense; bronchoconstriction; allergy; hyposecretion; pain; respiratory tract inflammation; adenosine sensitivity; lung; cancer;

analgesic; hypotensive; immunosupressive; cytostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory distress syndrome; allergic thinitis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis; pulmonary transplantation rejection; ss; primer. surfactant depletion; antiallergic; antiinflammatory; antiasthmatic;

Homo sapiens.

Pabalan J, Aquilar D;

Katz E,

i Y, Sandrasagra A, Ka Tang L, Shahabuddin S;

Li Y,

WO200285309-A2.

31-OCT-2002.

23-APR-2002; 2002WO-US013143

24-APR-2001; 2001US-0286036P.

(EPIG-) EPIGENESIS PHARM INC.

Pabalan J, Aguilar D; Katz E, Li Y, Sandrasagra A, Ka Tang L, Shahabuddin S; Miller S, Nyce JW,

WPI; 2003-093058/08.

S Pharmaceutical composition for treating asthma, has antisense oligonucleotide containing less percentage of adenosine, targeted tnucleic acids associated with lung airway or lung dysfunction, and bronchodilating agent

Claim 15; SEQ ID NO 3189; 763pp; English.

This invention describes a novel composition (a) a first active agent, comprising oligonuclectides, effective for alleviating comprising oligonuclectides, effective for alleviating comprising oligonuclectides, effective for alleviating reducing adenosine sensitivity, levels of adenosine (A) or (A) receptors, cuffactant depletion or hyposecretion, when administered to a mammal. The cligonuclectides are derived from a gene encoding or regulating configuration of a target polypeptide associated with lung airway or lung expression of a target polypeptide associated with lung airway or lung dystunction also describes a kit, that comprises: (a) a delivery device, in separate containers, (b) the oligonuclectides, (c) activity is a contraction for adding a carrier and for use of the kit. The composition of the invention has antiallergic, antiinflammatory, antiasthmatic, configuratory, lung or malignant disease. The administered composition comprises oligo and is administered to reduce the production or availability, or to increase the degradation of the target mRNA or to reduce the amount of target polypeptide present in the lungs. The composition, and/or bronchoconstriction and/or lung configuration, and/or surfactant hypoproduction are associated with a disease or condition such as pulmonary vasoconstriction, configuration rejection, pulmonary impeded respiration; repliratory configuration rejection, pulmonary impeded respiration; pulmonary configuration rejection, pulmonary impeded respiration; pulmonary configuration rejection, pulmonary interess syndrome, pain, cystic fibrosis, allergic rhimitis, pulmonary to thymidines present in the target RNA serves to prevent the breakdown of the right present in the target RNA serves to prevent the breakdown of thymidines present in the ranget RNA serves to prevent the breakdown of the pulse present in the reduced th thereby, to brain, heart, kidney, etc, tissue environment and unwanted effects due to it lung, brain, heart, prevent any

Sequence 20 BP; 10 A; 2 C; 7 G; 1 T; 0 U; 0 Other;

Gaps .; 0 0.8%; Score 14.2; DB 1; Length 20; 14.2%; Pred. No. 8.7e+02; ve 0; Mismatches 3; Indels Query Match
Best Local Similarity 84.2%;
Matches 16; Conservative

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1008 CGAGAGGGAGAGCTCAAG 1026

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1 CCAGAGAGAGAGATCAAG 19

Sequence 20 BP; 5 A; 3 C; 7 G; 5 T; 0 U; 0 Other;

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analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory distress syndrome; allergic rhinitis; pulmonary hyportension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis; pulmonary transplantation rejection; ss; primer.
                                                                                       respiratory tract inflammation, adenosine sensitivity, lung, cancer, surfactant depletion, antiallergic, antiinflammatory, antiasthmatic,
                                                                               antisense; bronchoconstriction; allergy; hyposecretion; pain;
                                                             Human transglutaminase-derived oligo SEQ ID 19.
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        ABD21007 standard; DNA; 20 BP.
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                                           29-JUL-2004 (first entry)
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Tang L,
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                          ABD21007;
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ABD21007
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Pabalan J, Aguilar Katz E,

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Pharmaceutical composition for treating asthma, has antisense oligonuclectide containing less percentage of adenosine, targeted tructeic acids associated with lung airway or lung dysfunction, and bronchodilating agent.

Claim 15; SEQ ID NO 19; 763pp; English.

This invention describes a novel composition (a) a first active agent, comprising oligonuclectides, effective for alleviating bronchoconstriction, respiratory tract inflammation, allergies and reducing adenosine sensitivity, levels of adenosine (A) or (A) receptors, surfactant depletion or hyposecretion, when administered to a mammal. The oligonucleotides are derived from a gene encoding or regulating expression of a target polypeptide associated with lung airway or lung dysfunction or cancer and can be anti-sense to the corresponding mRNA. The invention also describes a kit, that comprises: (a) a delivery instructions for adding a carrier and for use of the kit. The composition of the invention has antiallergic, antiinflammatory, antiasthmatic, inflammation, allergies, asthma, impeded respiration, respiratory distress syndrome, pain, cystic fibrosis, allergic rhinitis, pulmonary hypertension, emphysema, chronic obstructive pulmonary disease, pulmonary transplantation rejection, pulmonary infections, bronchitis or cancer. The reduced adenosine content of the anti-sense oligos corresponding to thymidines present in the target RNA serves to prevent the breakdown of the oligonuclectides into products that free adenosine into the system e.g., lung, brain, heart, kidney, etc, tissue environment and thereby, to prevent any unwanted effects due to it analgesic, hypotensive, immunosuppressive and cytostatic activity, is a beta-adrenergic agonist. The composition is useful for preventing or treating a respiratory, lung or malignant disease. The administered composition comprises oligo and is administered to reduce the production or availability, or to increase the degradation of the target mRNA or reduce the amount of target polypeptide present in the lungs. The pulmonary obstruction, and/or surfactant hypoproduction are associated inflammation, allergies and/or surfactant hypoproduction are associated with a disease or condition such as pulmonary vasoconstriction,

analgesic, hypotensive, immunosuppressive and cytostatic activity, is a beta-adrenergic agonist. The composition is useful for preventing or treating a respiratory, lung or malignant disease. The administered composition comprises oligo and is administered to reduce the production or availability, or to increase the degradation of the target mRNA or to reduce the amount of target polypeptide present in the lungs. The

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                                                                                                                                                                                                                                                                                                                                   Human; antisense; bronchoconstriction; allergy; hyposecretion; pain; respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depoletion; antiallergic; antiinflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory distress syndrome; allergic rhinitis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis; pulmonary transplantation rejection; ss; primer.
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                                           Gaps
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   Score 14.2; DB 1; Length 20;
Pred. No. 8.7e+02;
0; Mismatches 3; Indels
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Tang L, Shahabuddin
                                                                                                           19 GAACCIGCICAICTCCAAG 1
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   0.8%;
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                                                                                                                                                                                              ABD21288 standard; DNA; 20
                                                                                                                                                                                                                                                                   (first entry)
                                      Conservative
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                     Similarity
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Query Match
Best Local Simi:
Matches 16;
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Miller S,
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expression of a target polypeptide associated with lung airway or lung

The invention or cancer and can be anti-sense to the corresponding arguery dysfunction or cancer and can be anti-sense to the corresponding arguery and carcer and can be anti-sense; (a) a delivery device, in separate containers, (b) the oligonucleotides, (c) instructions for adding a carrier and for use of the kit. The composition of the invention has antiallergic, antiinflammatory, antiasthmatic, analgesic, bypotensive, immunosuppressive and cytostatic activity, is a beta-adrenergic agonist. The composition is useful for preventing or treating a respiratory, lung or malignant disease. The administered composition comprises oligo and is administered to reduce the production or availability, or to increase the degradation of the target mRNA or reduce the amount of target polypeptide present in the lungs. The reduce the amount of target polypeptide present in the lungs. The inflammation, allergies and/or surfactant hypoproduction are associated with a disease or condition, ench as pulmonary vasoconstriction, the anismation, allergies, asthma, impeded respiration, respiratory bypertension, emphysema, chronic obstructive pulmonary disease, pulmonary the reduced adenosine content of the anti-sense oligos or cancer. The reduced adenosine content of the anti-sense oligos corresponding to thymidines present in the target RNA serves to prevent the breakdown of thymidines present in the target RNA serves to prevent the breakdown of thymidines present in the target RNA serves to prevent the breakdown of the oligonucleonia.

e.g., lung, brain, heart, kidney, etc, tissue environment and thereby, to prevent any unwanted effects due to it

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inflammation, allergies and/or surfactant hypoproduction are associated with a disease or condition such as pulmonary vasoconstriction, inflammation, allergies and/or surfactant hypoproduction are spiratory distances or condition such as pulmonary vasoconstriction, respiratory distress syndrome, pain, cystic fibrosis, allergic rhintis, pulmonary hypertension, emphysema, chronic obstructive pulmonary disease, pulmonary Transplantation rejection, pulmonary infections, bronchitis or cancer. The reduced adenosine content of the anti-sense oliges corresponding to thymidines present in the target RNA serves to prevent the breakdown of the oligonucleotides into products that free adenosine into the system e.g., lung, brain, heart, kidney, etc, tissue environment and thereby, to prevent any unwanted effects due to it
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obstruction, and/or bronchoconstriction and/or lung
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Pred. No. 8.7e+02;
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Best Local Similarity 84.2
Matches 16; Conservative
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                                                                                                                                                                                                                                                                                                                 0.8%; Score 14.2; DB 1; Length 20;
84.2%; Pred. No. 8.7e+02;
iive 0; Mismatches 3; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human myosin X-derived oligonucleotide SEQ ID 2751.
                                                                                                                                                                                                                                                                                             Sequence 20 BP; 4 A; 6 C; 6 G; 4 T; 0 U; 0 Other;
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nes 16; Conservative
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Tang L,
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This invention describes a novel composition (a) a first active agent, comprising oligonuclectides, effective for alleviating bronchoconstriction, respiratory tract inflammation, allergies and reducing adenosine sensitivity, levels of adenosine (A) or (A) receptors, surfactant depletion or hyposecretion, when administered to a mammal. The oligonuclectides are derived from a gene encoding or regulating Claim 15; SEQ ID NO 509; 763pp; English

Pharmaceutical composition for treating asthma, has antisense oligonucleotide containing less percentage of adenosine, targeted trucleic acids associated with lung airway or lung dysfunction, and

bronchodilating agent.

Aguilar

Pabalan J,

Katz E,

Li Y, Sandrasagra A, Ka Tang L, Shahabuddin S;

Miller S,

Nyce JW,

WPI; 2003-093058/08

24-APR-2001; 2001US-0286036P (EPIG-) EPIGENESIS PHARM INC

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This invention describes a novel composition (a) a first active agent, comprising oligonucleotides, effective for alleviating bronchoconstriction, respiratory tract inflammation, allergies and reducing adenosine sensitivity, levels of adenosine (A) or (A) receptors, surfactant depletion or hyposecretion, when administered to a mammal. The colligonucleotides are derived from a gene encoding or regulating expression of a target polypeptide associated with lung airway or lung expression of a target polypeptide associated with lung airway or lung dysfunction or cancer and can be anti-sense to the corresponding mRNA. The invention also describes a kit, that comprises: (a) a delivery device, in separate containers, (b) the oligonucleotides, (c) instructions for adding a carrier and for use of the kit. The composition of the invention has antiallergic, antiinflammatory, antiasthmatic, analgesic, hypotensive, immunosuppressive and cytostatic activity, is a beta-adrenergic agonist. The composition is useful for preventing or treating a respiratory, lung or malignant disease. The administered composition comprises oligo and is administered to reduce the amount of target polypeptide present in the lungs. The pulmonary obstruction, and/or surfactant hypoproduction are associated inflammation, allergies and/or surfactant hypoproduction are associated with a disease or condition such as pulmonary vasconarication.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 inflammation, allergies, asthma, impeded respiration, respiratory distress syndrome, pain, cystic fibrosis, allergic rhinitis, pulmonary hypertension, emphysema, chronic obstructive pulmonary disease, pulmonary
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        the oligonucleotides into products that free adenosine into the system e.g., lung, brain, heart, kidney, etc, tissue environment and thereby, to prevent any unwanted effects due to it
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      transplantation rejection, pulmonary infections, bronchitis or cancer. The reduced adenosine content of the anti-sense oligos corresponding to thymidines present in the target RNA serves to prevent the breakdown of
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pituitary tumour-transforming gene 1; securin; TUTR1;
ESP-1 associated protein; cytostatic; antiinflammatory; inflammation;
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/note= "Phophorothioate backbone.All cytidines are 5-
methylcytidines"
oligonucleotide containing less percentage of adenosine, targeted to nucleic acids associated with lung airway or lung dysfunction, and
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84.2%; Pred. No. 8.7e+02;
ative 0; Mismatches 3; Indels
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                                                                                            Claim 15; SEQ ID NO 2751; 763pp; English.
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                                               pronchodilating agent.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Key
modified_base
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADG42473;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      tumour.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RESULT 1093
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADG42473,
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New antisense compound targeted to a nucleic acid molecule encoding human pituitary tumor-transforming gene useful for prophylaxis of e.g. inflammation and tumor formation.
                                                                                                                                                                                                                                                                                                                                                                                                                                    The invention relates to an antisense compound having a length of 8 - 50 nucleotides and targeted to a nucleic acid molecule encoding human pituliary tumour-transforming gene (PTG1, also known as securin, TUTR1 or ESP-1 associated protein). The antisense oligonucleotides are useful for inhibiting the expression of PTG1 in human cells or tissues, for treating diseases associated with PTTG1, as tools in differential and/or combinatorial analysis to elucidate expression patterns of a portion of the entire complement of genes expressed within cells and tissues, for diagnostics, therapeutics, and prophylaxis of e.g. inflammation, tumour formation and as research reagents and kits. The present sequence is an
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                antisense oligonucleotide of the invention, targeting human PTTG1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MUC-1; cytostatic; vaccine; tumour; carcinoma; immune response; cytotoxic T lymphocyte; antibody response; human; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0.8%; Score 14.2; DB 1; Length 20;
14.2%; Pred. No. 8.7e+02;
ve 0; Mismatches 3; Indels
/mod_base= OTHER
/note= "2'-methoxyethyl nucleotides"
16. .20
                                                                            /note= "2'-methoxyethyl nucleotides"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Seguence 20 BP; 6 A; 6 C; 2 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                            Example 15; SEQ ID NO 30; 46pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MUC-1 related PCR primer 2006MUC1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1010 AGAGGGGAGAGCTCAAGCT 1028
                                                                /mod base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         19 AGATGGGAGATCTCAAGTT 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       BP.
                                                                                                                                                                                                       02-APR-2002; 2002US-00114683.
                                                                                                                                                                          02-APR-2002; 2002US-00114683.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            84.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADE44005 standard; DNA; 20
                                                /*tag= c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             16; Conservative
                                                                                                                                                                                                                                     (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                                                                                 WPI; 2004-041251/04.
                                                                                                                                                                                                                                                     (ABBO ) ABBOTT LAB.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Similarity
                                                                                                                                                                                                                                                                                    Luo Y;
                                                                                                              US2003194396-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WO2003099193-A2,
                                modified base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens.
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                                                                                                                                           16-OCT-2003.
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                                                                                                                                                                                                                                                                                  Watt AT,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADE44005;
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Best Local S
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Burden N, Ellis JH, Hamblin PA;
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XX
AC ADF32644;
XX
AC ADF32644;
XX
DT 26-FEB-2004 (first entry)
XX
WUC-1 related PCR primer 2006MUC
XX
WWC-1 antigen; immune response; l
XW
WWRR: repeat unit; tumour; metas
XW
WOZO03100060-A2.
XX
XX
YX
WOZO03100060-A2.
XX
PP 04-DEC-2003.
XX
PP 23-MAY-2002; 2002GB-00012046.
XX
PR 24-MAY-2002; 2002GB-00012046.
XX
PR 24-MAY-2002; 2002GB-0012046.
XX
XX
PR Z4-MAY-2002; Z002GB-0012046.
XX
XX
PR Z4-MAY-2002; Z002GB-0011046.
XX
XX
XX
PR Z4-MAY-2003; Z003WO-EP005594.
XX
XX
YI Burden N, Ellis JH, Hamblin PA
                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity 84.29
----hes 16; Conservative
                                                                                                                                                                                                                                                                                                       the present invention.
                                                      WPI; 2004-035026/03.
                                     Burden N,
                                                                                                                                                                                                                                                                                                                                                                                                                               RESULT 1095
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The present invention describes a nucleic acid molecule which encodes a MUC-1 antigen. The nucleic acid is capable of raising an immune response in vivo, has reduced susceptibility to recombination than full-length MUC-1 and comprises between 1 and 15 variable number of tandem repeats (WNTR) perfect repeat units. Also described: (1) a plasmid comprising the DNA molecule; (2) a protein encoded by the nucleic acid, (3) a pharmaceutical composition comprising the nucleic acid, plasmid or protein and an excipient, dilunent or carrier; and (4) a method of treating or preventing tumours or metastases. A MUC1 antigen has cytostatic activity, and can be used in vaccines, and in gene therapy. The nucleic acid is useful for preparing a composition for treating or preventing tumours or metastases. The present sequence is used in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ss; E2-EPF; autoimmune disease; skin disease; pemphigus foliaceus; human;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New antisense oligonucleotide targeted to a nucleic acid encoding E2-EPF, useful for treating an autoimmune disease, preferably a skin disease, such as, pemphigus foliaceus.
                                                New nucleic acid molecule encoding a MUC-1 antigen, useful for preparing
                                                                       a composition for treating or preventing tumors or metastases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0.8%; Score 14.2; DB 1; Length 20;
84.2%; Pred. No. 8.7e+02;
tive 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        antisense oligonucleotide ISIS 156928.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 20 BP; 6 A; 9 C; 4 G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 exemplification of the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example 15; SEQ ID NO 39; 45pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1015 GGAGAGCTCAAGCTGGCTG 1033
                                                                                                                        Example; Page 66; 66pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 20 gdAgrecrerrecrescre 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         14-JUN-2002; 2002US-00173240.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADG72400 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Best Local Similarity 84.2
Matches 16, Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Dobie KW;
WPI; 2004-042811/04.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2004-052171/05.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human E2-EPF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Homo sapiens,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          11-MAR-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        18-DEC-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Monia BP,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     antisense.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADG72400;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        RESULT 1096
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ò
                                                                                                                                                                                                                                                                                                                                                           The present invention describes a nucleic acid molecule encoding a MUC-1 derivative that is devoid of all perfect repeats. Also described: (1) a plasmid comprising the DNA molecule; (2) a protein encoded by a nucleic acid molecule; (3) a pharmaceutical composition comprising the nucleic acid, the plasmid or the protein and a pharmaceutical acceptable excipient, diluent or carrier; and (4) a method of treating or preventing tuncheic acid, plasmid, a protein carrier and can be used in vaccines. The nucleic acid, plasmid, a protein or the pharmaceutical composition of the protein can be used in the preparation of a medicament for the treatment or prevention MUC-1 expressing tuncurs. The tunour can be carcinomas of the breast, lung, gastric or other gastrointestinal carcinomas. The tuncleic acid vaccines are easy to produce in large quantities compared over conventional protein vaccination. Seen at small doses they have been reported to induce strong immune responses and can induce a cytotoxic T lymphocyte immune response as well as an antibody response. The present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MUC-1 antigen; immune response; MUC-1; variable number of tandem repeat; VNTR; repeat unit; tumour; metastasis; cytostatic; vaccine; gene therapy;
                                                                                                                                                                                            New nucleic acid molecule encoding a MUC-1 derivative that is devoid of all perfect repeats, useful as vaccine for treating or preventing MUC-1 expressing tumors e.g. carcinoma of the breast, lung or gastrointestinal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 sequence represents a PCR primer which is used in the exemplification of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 14.2; DB 1; Length 20;
Pred. No. 8.7e+02;
0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 20 BP; 6 A; 9 C; 4 G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MUC-1 related PCR primer 2006MUC1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1015 GGAGAGCTCAAGCTGGCTG 1033
                                                                                                                                                                                                                                                                                                                       Example; Page 34; 34pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           20 géAgrécrerrecredere 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0.8%;
     24-MAY-2002; 2002GB-00012036
                                                   (GLAX ) GLAXO GROUP LTD
                                                                                                   Hamblin P;
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The invention relates to a compound targeted to, and which specifically hybridises with, a nucleic acid molecule encoding E2-EPF, and that inhibits the expression of E2-EPF. The compound, composition and methods are useful for treating a disease or condition associated with E2-EPF, such as an autoimmune disease, preferably a skin disease such as

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Gaps ., 10017621-381.rng

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ADG72433 RESULT

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The invention relates to a compound targeted to, and which specifically hybridises with, a nucleic acid molecule encoding E2-EPF, and that inhibits the expression of E2-EPF. The compound, composition and methods are useful for treating a disease or condition associated with E2-EPF, seuch as an autoimmune disease, preferably a skin disease such as pemphigus foliaceus. They are also useful in research and diagnostics for modulating the expression of E2-EPF. The present sequence represents a human E2-EPF antisense oligonucleotide.
                                                                                                                 E2-EPF; autoimmune disease; skin disease; pemphigus foliaceus; human;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New antisense oligonucleotide targeted to a nucleic acid encoding E2-EPF, useful for treating an autoimmune disease, preferably a skin disease, such as, pemphigus foliaceus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ss; E2-EPF; autoimmune disease; skin disease; pemphigus foliaceus; human.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0.8%; Score 14.2; DB 1; Length 20; 84.2%; Pred. No. 8.7e+02;
                                                                               antisense oligonucleotide ISIS 156921.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 20 BP; 2 A; 7 C; 4 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example 15; SEQ ID NO 32; 45pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             target region ISIS 72344.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1080 CAATGAGGTGGTGACACTG 1098
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          19 CAAGGAGGTGACGACACTG 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                BP.
                                                                                                                                                                                                                                                                                                            14-JUN-2002; 2002US-00173240.
                                                                                                                                                                                                                                                                                                                                                  14-JUN-2002; 2002US-00173240.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       14-JUN-2002; 2002US-00173240.
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                                         (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DNA;
                                                                                                                                                                                                                                                                                                                                                                                        (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                            Dobie KW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2004-052171/05
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADG72427 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Best Local Similarity
                                                                                                                                                                                                                                   US2003232436-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           US2003232436-A1.
                                                                                                                                                                                                 Homo sapiens.
                                                                               Human E2-EPF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human B2-EPF
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                                         11-MAR-2004
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                                                                                                                                       antisense.
                                                                                                                                                                                                                                                                                                                                                                                                                            Monia BP,
                                                                                                                                                                             Synthetic.
     ADG72393;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The invention relates to a compound targeted to, and which specifically hybridises with, a mucleic acid molecule encoding EO-EPF, and that inhibits the expression of E2-EPF. The compound, composition and methods are useful for treating a disease or condition associated with E2-EPF, such as an autoimmune disease, preferably a skin disease such as pemphigus foliaceus. They are also useful in research and diagnostics for modulating the expression of E2-EPF. The present sequence represents a human E2-EPF target region.
   diagnostics for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ss; E2-EPF; autoimmune disease; skin disease; pemphigus foliaceus; human.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New antisense oligonucleotide targeted to a nucleic acid encoding E2-EPF, useful for treating an autoimmune disease, preferably a skin disease, such as, pemphigus foliaceus.
pemphigus foliaceus. They are also useful in research and diagnostics f modulating the expression of E2-EPF. The present sequence represents a human E2-EPF antisense oligonucleotide.
                                                                                                                                                         Gaps
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0
                                                                                                               Query Match 0.8%; Score 14.2; DB 1; Length 20; Best Local Similarity 84.2%; Pred. No. 8.7e+02; Matches 16; Conservative 0; Mismatches 3; Indels
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84.2%; Pred. No. 8.7e+02;
trive 0; Mismatches 3; Indels
                                                                           Sequence 20 BP; 5 A; 5 C; 6 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 20 BP; 4 A; 6 C; 5 G; 5 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                              Human E2-EPF target region ISIS 72351.
                                                                                                                                                                                             CTGCTCCTGGGGAACTTCG 293
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       275 CTGCTCCTGGGGAACTTCG 293
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es 16; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2004-052171/05.
                                                                                                                                                                                                                                                                                                                            ADG72433 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             US2003232436-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Homo sapiens
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ADG72393/c
ID ADG7239:
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The invention relates to an antisense oligonucleotide comprising 8-80 nucleobases in length targeted to the coding region of a nucleic acid molecule encoding PPAR-delta (peroxisome proliferative activated receptor delta), where the antisense compound inhibits the expression of the PPAR-delta and has any of the 66 sequences of 20 amino acids fully defined in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Example 16; SEQ ID NO 262; 155pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           31-MAY-2002; 2002US-00160807.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             31-MAY-2002; 2002US-00160807.
                                                                                                                                                                                                                                                                                                              ADG87026 standard; cDNA; 20
                                                                                                                                                                            numan B2-EPF target region.
                                                                                                                                                                                                                       Local Similarity 84.2
nes 16; Conservative
       (ISIS-) ISIS PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (ISIS-) ISIS PHARM INC
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                         Monia BP, Dobie KW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2004-022078/02.
                                         WPI; 2004-052171/05.
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the specification. Also included are a compound of 8-80 nucleobases in length that specifically hybridises with at least an 8-nucleobase portion of a preferred target region on a nucleic acid molecule encoding PpAR-delta and a composition comprising the antisense oligonucleotide and a carrier. The antisense oligonucleotide comprises at least one modified internucleoside linkage (preferably a phosphorothicate linkage), at least one sugar moiety (preferably 2'-O-methoxyethyl moiety) and at least one modified nucleobase (which is a 5-methyl cytosine). The antisense compounds are useful for treating cancer, osteoporosis, diabetes or various endocrine disorders. The Human PPAR delta gene is located on sequence for the antisense oligonucleotides of the invention. Mouse, ss, PPAR delta; peroxisome proliferative activated receptor delta; antisense gene therapy; cytostatic; osteopathic; antidiabetic; cancer; osteoporosis; diabetes; endocrine disorder. 1. .20 /*tag= b//mod_bas= OTHER /note= "Phosphorothioate linkages and all cytidines are -methylcytidines" 0.8%; Score 14.2; DB 1; Length 20; 34.2%; Pred. No. 8.7e+02; Sequence 20 BP; 3 A; 3 C; 9 G; 5 T; 0 U; 0 Other; Mouse PPAR antisense oligonucleotide ISIS 221095. /note= "2'-methoxyethyl residue"
16. .20 /note= "2'-methoxyethyl residue" 0; Mismatches Location/Qualifiers /*tag= c /mod_base= OTHER 'mod base= OTHER 344 TGAAGATGGGGTCTGATGG 362 2 recadaredecrerentes 20 BP 31-MAY-2002; 2002US-00160807. 31-MAY-2002; 2002US-00160807. 84.2%; ADG86888 standard; DNA; 20 (first entry) 16; Conservative *tag= (ISIS-) ISIS PHARM INC Local Similarity US2003224514-A1 modified base modified base modified base Mus musculus 11-MAR-2004 04-DEC-2003 ADG86888 Query Match Matches ADG86888, 88888888888888888 q ð . 0 Mouse; ss; PPAR delta; peroxisome proliferative activated receptor delta; antisense gene therapy; cytostatic; osteopathic; antidiabetic; cancer; osteoporosis; diabetes; endocrine disorder. The invention relates to a compound targeted to, and which specifically hybridises with, a nucleic acid molecule encoding E2-EPF, and that inhibits the expression of E2-EPF. The compound, composition and methods are useful for treating a disease or condition associated with E2-EPF, such as an autolimmune disease, preferably a skin disease such as pemphigus foliaceus. They are also useful in research and diagnostics for modularing the expression of E2-EPF. The present sequence represents a New antisense oligonucleotide targeted to a nucleic acid encoding E2-EPF, useful for treating an autoimmune disease, preferably a skin disease, such as, pemphigus foliaceus. New antisense oligonucleotides of 8-80 nucleobases, useful for treating cancer, diabetes, osteoporosis or various endocrine disorders. Gaps ; 0.8%; Score 14.2; DB 1; Length 20; 84.2%; Pred. No. 8.7e+02; iive 0; Mismatches 3; Indels Mouse PPAR antisense oligonucleotoide target sequence #22. Sequence 20 BP; 7 A; 4 C; 7 G; 2 T; 0 U; 0 Other; Example 15; SEQ ID NO 66; 45pp; English. 1080 CAATGAGGTGGTGACACTG 1098 2 chaggaggrgacgachcrg 20 Watt AT ВР (first entry)

New antisense oligonucleotides of 8-80 nucleobases, useful for treating cancer, diabetes, osteoporosis or various endocrine disorders.

Watt AT;

Gaarde W, Freier SM, WPI; 2004-022078/02. Example 16; SEQ ID NO 124; 155pp; English

10017621-381.rng

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The invention relates to an antisense oligonucleotide comprising 8-80 nucleobases in length targeted to the coding region of a nucleic acid molecule encoding PPAR-delta (peroxisome proliferative activated receptor delta), where the antisense compound inhibits the expression of the PPAR-delta delta and has any of the 66 sequences of 20 amino acids fully defined in the specification. Also included are a compound of 8-80 nucleobases in length that specifically hybridises with at least an 8-nucleobase portion of a preferred target region on a nucleic acid molecule encoding PPAR-delta and a composition comprising the antisense oligonucleotide and a carier. The antisense oligonucleotide comprises at least one modified internucleoside linkage (preferably 2'-O-methoxyethyl moiety) and at least one modified nucleobase (which is a 5-methyl cytosine). The antisense compounds are useful for treating cancer, osteoporosis, diabetes or various endocrine disorders. The Human PPAR delta gene is located on chromosome 6p21. The present sequence is an antisense oligonucleotide of
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The invention relates to a novel antisense compound targeted to a nucleic acid molecule encoding human apolipoprotein B (ApoB) which specifically hybridises with and inhibits the expression of human apolipoprotein B. The compound of the invention demonstrates antiarteriosclerotic,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    anorectic, lipid; cholesterol metabolism; atherosclerosis; diabetes Type 2; obesity; hyperlipidaemia; cardiovascular; gene therapy; antisense; 2'-0-methoxyethyl gapmer; phosphorothioate backbone; 2'-MOE;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New antisense compound, useful for preparing a composition for treating abnormal lipid or cholesterol metabolism, atherosclerosis, diabetes Type 2, obesity, hyperlipidemia or cardiovascular disease.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                          0.8%; Score 14.2; DB 1; Length 20; 84.2%; Pred. No. 8.7e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                3; Indels
                                                                                                                                                                                                                                                                                                                                                                   Sequence 20 BP; 5 A; 9 C; 3 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                0; Mismatches
                                                                                                                                                                                                                                                                                                                                   the invention targeting mouse PPAR delta.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         344 TGAAGATGGGGTCTGATGG 362
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             BP,
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13-NOV-2002; 2002US-0426324P
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                                                                                                                                                                                                                                                                                                                                                                                                                              Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 human; ss.
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cardiant, antidiabetic and anorectic activities and may be useful for preparing a composition for treating abnormal lipid or cholesterol metabolism, atherosclerosis, diabetes Type 2, obesity, hyperlipidaemia or cardiovascular disease. Furthermore, the compound has gene therapy applications. The current sequence is that of the 2'-O-methoxychthyl (2'-MOE) gapmer antisense oligo of the invention which has 2'-MOE 'wings', a phosphorothioate backbone throughout and in which all cytidine residues
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       acid molecule encoding human apolipoprotein B (ApoB) which specifically hybridises with and inhibits the expression of human apolipoprotein B. The compound of the invention demonstrates antiarteriosclerotic, cardiant, antidiabetic and anorectic activities and may be useful for preparing a composition for treating abnormal lipid or cholesterol metabolism, atherosclerosis, diabetes Type 2, obesity, hyperlipidaemia or cardiovascular disease. Furthermore, the compound has gene therapy applications. The current sequence is that of the human ApoB antisense inhibition target DNA of the invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               apolipoprotein B; ApoB; antiarteriosclerotic; cardiant; antidiabetic; anorectic; lipid; cholesterol metabolism; atherosclerosis; diabetes Type 2; obesity; hyperlipidaemia; cardiovascular; gene therapy; antisense inhibition target; human; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New antisense compound, useful for preparing a composition for treating abnormal lipid or cholesterol metabolism, atherosclerosis, diabetes Type 2, obesity, hyperlipidemia or cardiovascular disease.
                                                                                                                                                                                                                                                       Gaps
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                                                                                                                                                                                                                                                       0
                                                                                                                                                                                                                 Length 20;
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                                                                                                                                                                                                               Score 14.2; DB 1; Length 2
Pred. No. 8.7e+02;
0; Mismatches 3; Indels
                                                                                                                                                                           Sequence 20 BP; 9 A; 5 C; 4 G; 2 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                     0; Mismatches
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                                                                                                                                                                                                                                                                                          583 CTATCTGAGATTGGCTTTG 601
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13-NOV-2002; 2002US-0426324P.
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                                                                                                                                                                                                                                                       16; Conservative
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                                                                                                                                       are 5-methylcytidines.
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                                                                                                                                                                                                                                   Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 for treating
diabetes Type
                                                                                                                                                                                                                                                                                                                                                                                         apolipoprotein B; ApoB; antiarteriosclerotic; cardiant; antidiabetic; anorectic; lipid; cholesterol metabolism; atherosclerosis; diabetes Type 2; obesity; hyperlipidaemia; cardiovascular; gene therapy; antisense; 2'-O-methoxyethyl gapmer; phosphorothioate backbone; 2'-MOB;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       phosphorothicate backbone throughout and in which all cytidine residues
                                                                                                                                                                                                                                                                                                                                              2'-MOE gapmer antisense oligo targeted to human ApoB DNA 1 - SEQ ID 128
Gaps
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  Indels
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  Mismatches
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                                             1012 AGGGGAGACTCAAGCTGG 1030
  ٠:
                                                                                         AGGTATGAGCTCAAGCTGG 20
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13-NOV-2002; 2002US-0426324P.
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                                                                                                                                                                                                       ADH18139 standard; DNA; 20
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  Conservative
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Best Local Similarity
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  16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Crooke RM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         human; ss.
                                                                                                                                                                                                                                                       ADH18139;
  Matches
                                                                                                                                                            RESULT 111
ADD118139/ADD18139/ADD118139/ADD118139/ADD118139/ADD118139/ADD118139/ADD118139/ADD118139/ADD118139/ADD118139/ADD118139/ADD118139/ADD118139/ADD118139/ADD118139/ADD118139/ADD118139/ADD118139/ADD118139/ADD118139/ADD118139/ADD118139/ADD118139/ADD118139/ADD118139/ADD118139/ADD118139/ADD118139/ADD118139/ADD118139/ADD118139/ADD118139/ADD118
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31-MAY-2002; 2002US-00159856. 31-MAY-2002; 2002US-00159856.

JS2003228689-A1

11-DEC-2003

(ISIS-) ISIS PHARM INC.

Dobie KW;

Freier SM,

/note= "This oligonucleotide has a phosphorothioate backbone and 2'-methyoxyethyl (2'-MOS) wings at the 5' and 3' ends, which are 5 nucleotides in length at each end. All cytidine residues are 5-methyloytidines.

Antisense therapy, human, G protein-coupled receptor kinase 6; GPCR kinase 6; GRK6; rheumatoid arthritis, drug addiction, uterine contractility; hypertension; aberrant haematopoiesis; antiinflammatory; antiarthritic; antirheumatic; hypotensive;

phosphorothioate; ss.

Homo sapiens

Location/Qualifiers /*tag= a /mod_base= OTHER

modified base

Human GRK6 DNA, antisense oligonucleotide #14.

(first entry)

25-MAR-2004

ADH48222;

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·:
                                                                                                                                                                                                                                                                                                                                                        The present invention relates to antisense compounds targeted to a nucleic acid encoding G protein-coupled receptor (GPCR) kinase 6 (GRK6). The antisense compound comprises an antisense oligonucleotide that specifically hybridises with the nucleic acid and inhibits the expression of GRK6. The antisense oligonucleotide is a chimeric oligonucleotide. The antisense oligonucleotide comprises at least one modified internucleoside linkage, preferably a phosphorothioate linkage. It also comprises at least one modified sugar moiety, preferably a 2'-0-methoxyethyl (2'-MOE) sugar moiety. The antisense oligonucleotide further comprises at least one modified nucleobase, preferably a 5'-methylcytosine. The antisense oligonucleotides are useful for the treatment of diseases such as rheumatoid arthritis, drug addiction, uterine contractility, hypertension, and diseases or conditions arising from aberrant
                                                                                New compounds, particularly antisense oligonucleotides targeted to a nucleic acid encoding G protein-coupled receptor (GPCR) kinase 6, useful for treating diabetes, drug addiction, uterine contractility and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              oligonucleotide used in the examples of the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The present sequence represents an antisense
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0.8%; Score 14.2; DB 1; Length 20; 84.2%; Pred. No. 8.7e+02; ive 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 20 BP; 5 A; 3 C; 6 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                        Example 15; SEQ ID NO 24; 58pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      843 TGAGTACCTGGACAAGGAC 861
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TGAGTTCCTGGAAAAGGTC 19
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WPI; 2004-052027/05.
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                                                                                                                                                                                                            hypertension
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Gaps

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3; Indels

0; Mismatches

1012 AGGGGAGAGCTCAAGCTGG 1030

16; Conservative

Matches

AGGTATGAGCTCAAGCTGG 1

19

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ADH48222 standard; DNA; 20 BP

RESULT 1105

ADH48222 ID ADH

RESULT 1106

10017621-381.rng

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WO2003099215-A2
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                                  04-DEC-2003
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                                                                                                                             SD,
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                                                                                                                             Crosby
                                                                                                                                                                                                                                                                                                                                                                                                             Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                 to nucleic acids encoding a mammalian glucocorticoid receptor. The antisense oligonucleotides of the invention are useful for preventing or delaying infection, inflammation or tumour formation. The antisense oligonucleotides are also useful for treating diabetes, obesity, cardiovascular disorders, hyperlipidaemia or Cushing's syndrome. The present DNA sequence represents an antisense oligonucleotide that targets the human glucocorticoid receptor gene NOTE: The present sequence contains 2'-methoxyethyl (2'-MOE) wings and a phosphorothioate backbone.
                                                                                                                                                                                                                                                                                                                                                  New antisense compound targeted to a nucleic acid molecule encoding mammalian glucocorticoid receptor, useful for treating diabetes, obesity, cardiovascular disorder, hyperlipidemia or Cushing's syndrome.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     antisense oligonucleotide; glucocorticoid receptor; infection; inflammation; tumour formation; diabetees; obesity; cardiovascular disorder; hyperlipidaemia; Cushing's syndrome; human; ss; phosphorothioate backbone; 2 - methoxyethyl; 2 '-MOE.
                                                                                                                             's syndrome; human; ss;
                                                                                                                                                                                                                                                                                                                                                                                                                        The invention comprises an antisense oligonucleotides that are targeted
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human glucocorticoid receptor-specific antisense oligonucleotide #1997.
                                                                               Human glucocorticoid receptor-specific antisense oligonucleotide #138
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                                                                                                      oligonucleotide; glucocorticoid receptor; infection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0.8%; Score 14.2; DB 1; Length 20; 84.2%; Pred. No. 8.7e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Indels
                                                                                                                 inflammation, tumour formation, diabetes, obesity, cardiovascular disorder, hyperlipidaemia, Cushing's phosphorothioate backbone, 2'-methoxyethyl, 2'-MOE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 20 BP; 2 A; 6 C; 3 G; 9 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                Claim 4; SEQ ID NO 138; 985pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            621 TAAGCTGGACAAACTGGGC 639
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  20 TAAGCAAGAGAACTGGGC 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             BP.
          BP.
                                                                                                                                                                                                                                  20-MAY-2003; 2003WO-US016084
                                                                                                                                                                                                                                                         20-MAY-2002; 2002US-0381857P
         ADH63304 standard; DNA; 20
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                                                         (first entry)
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Best Local Similarity 84.2
Matches 16, Conservative
                                                                                                                                                                                                                                                                                                      Nalseth AE;
                                                                                                                                                                                                                                                                               (PHAA ) PHARMACIA CORP.
                                                                                                                                                                                                                                                                                                                            WPI; 2004-035034/03.
                                                                                                                                                                                      WO2003099215-A2.
                                                                                                                                                               Homo sapiens.
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                                                        25-MAR-2004
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                                                                                                                                                                                                                                                                                                        Crosby SD,
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                                                                                                       antisense
                                  ADH63304;
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 ADH63304/c
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to nucleic acids encoding a mammalian glucocorticoid receptor. The antiense oligonucleotides of the invention are useful for preventing or delaying infection, inflammation or tumour formation. The antisense oligonucleotides are also useful for treating diabetes, obesity, cardiovascular disorders, hyperlipidaemia or Cushing's syndrome. The present DNA sequence represents an antisense oligonucleotide that targets the human glucocorticoid receptor gene. NOTE: The present sequence contains 2'-methoxyethyl (2'-MOE) wings and a phosphorothioate backbone.
                                                                                                                                                                                                                                                                                                                                                             New antisense compound targeted to a nucleic acid molecule encoding mammalian glucocorticoid receptor, useful for treating diabetes, obesity, cardiovascular disorder, hyperlipidemia or Cushing's syndrome.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         antisense oligonucleotide; glucocorticoid receptor; infection; inflammation; tumour formation; diabetes; obesity; cardious formation; hyperlipidemia; Cushing's syndrome; human; ss; phosphorothicate backbone; 2-methoxyethyl; 2.-MOE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The invention comprises an antisense oligonucleotides that are targeted
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human glucocorticoid receptor-specific antisense oligonucleotide #4237
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
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Pred. No. 8.7e+02;
0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 20 BP; 3 A; 5 C; 2 G; 10 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 4; SEQ ID NO 1997; 985pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            619 ATTAAGCTGGACAAACTGG 637
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20-MAY-2003; 2003WO-US016084
                                                                        20-MAY-2002; 2002US-0381857P
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ADH67403 standard; DNA; 20
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les 16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Nalseth AE;
                                                                                                                                            (PHAA ) PHARMACIA CORP
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                                                                                                                                                                                                                                                                                             WPI; 2004-035034/03.
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                                                                                                                                            to nucleic acids encoding a mammalian glucocorticoid receptor. The antisense oligonucleotides of the invention are useful for preventing or delaying infection, inflammation or tumour formation. The antisense oligonucleotides are also useful for treating diabetes, obesity, present DNs sequence represents an antisense oligonucleotide that targets the human glucocorticoid receptor gene. NOTE: The present sequence contains 2'-methoxyethyl (2'-MOE) wings and a phosphorothioate backbone.
mammalian glucocorticoid receptor, useful for treating diabetes, obesity, cardiovascular disorder, hyperlipidemia or Cushing's syndrome.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New antisense compound targeted to a nucleic acid molecule encoding mammalian glucocorticoid receptor, useful for treating diabetes, obesity, cardiovascular disorder, hyperlipidemia or Cushing's syndrome.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            antisense oligonucleotide; glucocorticoid receptor; infection; inflammation; tumour formation, diabetee; obesity; cardiovascular disorder; hyperlipidaemia; Cushing's syndrome; human; ss; phosphorothioate backbone; 2 - methoxyethyl; 2'-MOE.
                                                                                                                               The invention comprises an antisense oligonucleotides that are targeted
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human glucocorticoid receptor-specific antisense oligonucleotide #4079
                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
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0
                                                                                                                                                                                                                                                                                                                                                                             Query Match 0.8%; Score 14.2; DB 1; Length 20; Best Local Similarity 84.2%; Pred. No. 8.7e+02; Matches 16; Conservative 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                        Sequence 20 BP; 5 A; 6 C; 6 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 4; SEQ ID NO 4079; 985pp; English.
                                                                                      Claim 4; SEQ ID NO 4237; 985pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  49 CCAGCAGTGTGACTGCTGA 67
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      20 ccaecaererecrrecrea 2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (PHAA ) PHARMACIA CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2004-035034/03.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADH67245;
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                                                                           Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New compounds, particularly antisense oligonucleotides targeted to a nucleic acid encoding vascular endothelial growth factor-C (WEGF-C), useful for treating atherosclerosis, diabetic retinopathy, or inflammatory disorders.
                                                                                                                                                                                                                                                                                                                                           human; ss; PCR; VEGF-C; cardiovascular disorder; atherosclerosis; diabetic retinopathy; autoimmune disorder; inflammatory disorder; vascular endothelial growth factor; primer.
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0
                                     Score 14.2; DB 1; Length 20;
Pred. No. 8.7e+02;
0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0.8%; Score 14.2; DB 1; Length 20;
84.2%; Pred. No. 8.7e+02;
artive 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Seguence 20 BP; 7 A; 6 C; 5 G; 2 T; 0 U; 0 Other;
             Sequence 20 BP; 5 A; 5 C; 7 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example 13; SEQ ID NO 5; 83pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1553 GGTCTTCGTCGATGCCTGA 1571
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                                                                                                                                    19 ccagcagrardcrigcrca 1
                                                                                                                                                                                                                   BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       17-JUN-2002; 2002US-00173718.
                                          0.8%;
                                                                                                         49 CCAGCAGTGTGACTGCTGA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    17-JUN-2002; 2002US-00173718.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  19 GGTCTTGTTCGCTGCCTGA
                                                                                                                                                                                                                                                                                                               Human VEGF-C PCR primer #1.
                                                                                                                                                                                                                   ADH54704 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADH50654 standard; DNA; 20
                                                                                                                                                                                                                                                                                (first entry)
                                                                         Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2004-061284/06.
                                         Query Match
Best Local Similarity
Matches 16; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Zhang H, Dobie KW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Local Similarity
les 16; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                          US2003232437-A1
                                                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens.
                                                                                                                                                                                                                                                                                25-MAR-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        18-DEC-2003.
                                                                                                                                                                                                                                                  ADH54704;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
                                                                                                                                                                                      RESULT 1110
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RESULT 1111
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Matches
                                                                                                                                                                                                      ADH54704/
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ID ADHS
XX SS
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Page

ADH50654;

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The present invention relates to antisense compounds targeted to a nucleic acid encoding interleukin-1 (IL-1) receptor-associated kinase-1 (IRAK-1). The antisense compound comprises an antisense oligonucleotide that specifically hybridises with the nucleic acid and inhibits the expression of IRAK-1. The antisense oligonucleotide is a chimeric oligonucleotide. The antisense oligonucleotide comprises at least one modified internucleoside linkage, preferably a phosphorothicate linkage. It also comprises at least one modified sugar moiety, preferably a 2-0-methoxyethyl (2-MOE) sugar moiety. The antisense oligonucleotide further comprises at least one modified antisense oligonucleotide further comprises at least one modified outleobase, preferably a 5-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  methylcytosine. The antisense oligonucleotides are useful for the treatment of diseases such as hyperproliferative disorders, e.g. cancer, autoimmune disorders, altered bone metabolism, and inflammation. The present sequence represents a human IRAK-1 DNA target sequence for an
                                                                                                                                                               Antisense therapy, human, interleukin-1 receptor-associated kinase-1, IL-1 receptor-associated kinase-1, IRAK-1, hyperprofilerative disorder e.g.; cancer; autoimmune disorder; altered bone metabolism or inflammation; cytostatic; immunosuppressive; osteopathic; antiinflammatory; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New compound having a sequence targeted to a nucleic acid encoding IL-1 receptor-associated kinase-1, useful for preparing a composition for treating hyperproliferative or autoimmune disorder or inflammation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example 15; SEQ ID NO 127; 66pp; English.
                                                                                                                        DNA target sequence #42.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Dobie KW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            10-JUN-2002; 2002US-00167034.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           10-JUN-2002; 2002US-00167034
                                                                         (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             antisense oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Freier SM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2004-052028/05.
                                                                                                                                                                                                                                                                                                                                                                           US2003228690-A1
                                                                                                                        Human IRAK-1
                                                                                                                                                                                                                                                                                                                              Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         25-MAR-2004
                                                                         25-MAR-2004
                                                                                                                                                                                                                                                                                                                                                                                                                             11-DEC-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Baker BF,
                       ADH50720;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADH61956;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1113
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADH61956
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    RESULT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ð
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        유
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ó:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The present invention relates to antisense compounds targeted to a nucleic acid encoding interleukin-1 (IL-1) receptor-associated kinase-1 (IRAK-1). The antisense compound comprises an antisense oligonucleotide that specifically hybridises with the nucleic acid and inhibits the expression of IRAK-1. The antisense oligonucleotide is a chimeric oligonucleotide. The antisense oligonucleotide comprises at least one modified sugar moiety, preferably a 2'-0- It also comprises at least one modified sugar moiety, preferably a 2'-0- methoxyethyl (2'-MOE) sugar moiety. The antisense oligonucleotide further methotycosine. The antisense oligonucleotide further methylotycosine. The antisense oligonucleotide further methylotycosine. The antisense oligonucleotide are useful for the treatment of diseases such as hyperproliferative disorders, e.g. cancer, treatment of diseases such as hyperproliferative disorders, e.g. cancer,
                                                                                                                                          Antisense therapy, human, interleukin-1 receptor-associated kinase-1, IL-1 receptor-associated kinase-1, IRAK-1; hyperproliferative disorder e.g., cancer; autoimmune disorder; altered bone metabolism or inflammation; cytostatic; immunosuppressive; osteopathic; antiinflammatory; phosphorothioate; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New compound having a sequence targeted to a nucleic acid encoding IL-1 receptor-associated kinase-1, useful for preparing a composition for treating hyperproliferative or autoimmune disorder or inflammation.
                                                                                                                                                                                                                                                                                                                                                                                                                       /mod_base= OTHER
/note= "This oligonucleotide has a phosphorothicate
/note= "This oligonucleotide has a phosphorothicate
backbone and 2'-methyoxyethyl (2'-MOE) wings at the 5'
and 3' ends, which are 5 nucleotides in length at each
end. All cytidine residues are 5-methylcytidines"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     autoimmune disorders, altered bone metabolism, and inflammation. The present sequence represents an antisense oligonucleotide used in the examples of the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0.8%; Score 14.2; DB 1; Length 20; 84.2%; Pred. No. 8.7e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 20 BP; 1 A; 7 C; 7 G; 5 T; 0 U; 0 Other;
                                                                                                DNA, antisense oligonucleotide #48.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Example 15; SEQ ID NO 61; 66pp; English.
                                                                                                                                                                                                                                                                                                                                                 Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             928 CAGCTGCTCCGTGGCCTGG 946
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Dobie KW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    10-JUN-2002; 2002US-00167034
                                                                                                                                                                                                                                                                                                                                                                                1. .2.
/*tag= a
                                                 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match 0.8
Best Local Similarity 84.2
Matches 16; Conservative
                                                                                                                                                                                                                                                                                                                                                                           . 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Freier SM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2004-052028/05.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               US2003228690-A1
                                                                                                                                                                                                                                                                                                                                                      Key
modified_base
                                                                                                   Human IRAK-1
                                                                                                                                                                                                                                                                                                    Homo sapiens
                                                 25-MAR-2004
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Baker BF,

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0
                                                                                 Gaps
                                                                                 ;
                                     0.8%; Score 14.2; DB 1; Length 20; 44.2%; Pred. No. 8.7e+02;
                                                                                 Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                            Basidiomycete; dioxin; ribosomal DNA; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                    Panellus stypticus rDNA PCR primer ITS 2, SEQ ID 2.
Sequence 20 BP; 5 A; 7 C; 7 G; 1 T; 0 U; 0 Other;
                                                                                 0; Mismatches
                                                                                                                          928 CAGCTGCTCCGTGGCCTGG 946
                                                                                                                                                                cascretrerecrectes
                                                                                                                                                                                                                                                                           BP.
                                                           84.2%;
                                                                                                                                                                                                                                                                         ADH61956 standard; DNA; 20
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ADH50720 standard; DNA; 20

RESULT 1112 ADH50720/c

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01-MAR-2002; 2002JP-00055681.
                                                                                                                                                                                                                                                                             01-MAR-2002; 2002JP-00055681
                                                                   dioxin degrading activity.
                                                                                                                                                                 Query Match 0.8
Best Local Similarity 84.2
Matches 16; Conservative
                                                      WPI; 2004-172258/17
   Panellus stypticus
                                                                                                                                                                                                                                                     Panellus stypticus
           JP2003250520-A.
                                                                                                                                                                                                                                                             JP2003250520-A
                                                                                                                                                                                                                             25-MAR-2004
                   09-SEP-2003
                                                                                                                                                                                                                                                                     09-SEP-2003
                                                                                                                                                   invention.
                                                                                                                                                                                                                      ADH61957;
                                                                                                                                                                                                      RESULT 1114
                                                                                                                                                                                                          ADH61957/
à
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the initial stage concentration of the first culture medium. A reagent destroying microbial cells is added to the culture medium. A reagent culture completion. The microbial cells are lysed and the residual dioxin concentration of both the culture medium are compared. The basidiomycete having the dioxin degrading activity is selected by screening, which involves selecting the microorganisms in which the residual dioxin concentration of the first culture medium is less than the residual dioxin concentration of the second culture medium. ADH61959-ADH61961 are tibosomal DNAs from Panellus stypticus used to determine the genus of the basidiomycetes used in the method of the invention. ADH61955-ADH61958 are PCR primers used to amplify the rDNA sequences in an example from the using basidiomycete such as Panellus stypticus having using decomposable basidiomycetes (Panellus stypticus). In (M1), the basidiomycete is cultured in a first culture medium containing dioxin and in a second culture medium not containing dioxin. The dioxin is added to the second culture medium such that its concentration becomes equal to cytostatic; cardiant; antiinflammatory; antimicrobial; antisense therapy; bhosphodiesterase 4D; cancer; cardiovascular disease; inflammation; inflection; inflammation; tumour formation; antisense technology; human; ss. The present invenion relates to a method (M1) for decomposing dioxin by /mod_base= OTHER /note= "OTHER= Phosphorothioate backbone. All cytidines Score 14.2; DB 1; Length 20; Pred. No. 8.7e+02; 0; Mismatches 3; Indels /mod_base= OTHER /note= "OTHER= 2'-0-Methoxyethyl (2'-MOE) 15. .20 Human phosphodiesterase 4D antisense oligonucleotide #18. Sequence 20 BP; 7 A; 5 C; 6 G; 2 T; 0 U; 0 Other; (DOKU-) DOKURITSU GYOSEI HOJIN SANGYO GIJUTSU SO. Example 3; SEQ ID NO 3; 11pp; Japanese. are 5-methylcytidines' Location/Qualifiers 1549 CTTCGGTCTTCGTCGATGC 1567 19 CTGCGTTCTTCATCGATGC 1 BP 0.8%; ADI15592 standard; DNA; 20 (first entry) Degrading dioxin by using dioxin degrading activity. Д /*tag= a Query Match 0.8 Best Local Similarity 84.2 Matches 16; Conservative 1. .20 /*tag= WPI; 2004-172258/17. Key modified_base modified base modified base 22-APR-2004 Homo sapiens ADI15592; RESULT 1115 AD115592 g ð ; The present invenion relates to a method (MI) for decomposing dioxin by using decomposable basidiomycetes (Panellus stypticus). In (MI), the basidiomycete is cultured in a first culture medium containing dioxin and in a second culture medium not containing dioxin. The dioxin is added to the second culture medium such that its concentration becomes equal to the initial-stage concentration of the first culture medium. A reagent destroying microbial cells is added to the culture medium after the culture completion. The microbial cells are lysed and the residual dioxin concentration of both the culture mediums are compared. The basidiomycete having the dioxin degrading activity is selected by screening, which involves selecting the microorganisms in which the residual dioxin concentration of the first culture medium is less than the residual dioxin dioxin concentration of the second culture medium. ADHG195-ADHG1961 Degrading dioxin by using basidiomycete such as Panellus stypticus having ribosomal DNAs from Panellus stypticus used to determine the genus of the basidiomycetes used in the method of the invention. ADH61955-ADH61958 are PCR primers used to amplify the rDNA sequences in an example from the Gaps ö 0.8%; Score 14.2; DB 1; Length 20; 84.2%; Pred. No. 8.7e+02; ive 0; Mismatches 3; Indels Basidiomycete; dioxin; ribosomal DNA; PCR; primer; ss. Panellus stypticus rDNA PCR primer ITS 3, SEQ ID 3. Sequence 20 BP; 2 A; 6 C; 5 G; 7 T; 0 U; 0 Other; (SAOC) MERCIAN CORP. (DOKU-) DOKURITSU GYOSEI HOJIN SANGYO GIJUTSU SO. Example 3; SEQ ID NO 2; 11pp; Japanese. 1549 CTTCGGTCTTCGTCGATGC 1567 2 crecerretrearcearee 20 ADH61957 standard; DNA; 20 BP 01-MAR-2002; 2002JP-00055681. 01-MAR-2002; 2002JP-00055681 (first entry)

ö

Gaps

. 0

nucleotides"

nucleotides"

/note= "OTHER= 2'-O-Methoxyethyl (2'-MOE)

base= OTHER

JS2003220273-A1

27-NOV-2003

(SAOC) MERCIAN CORP

/*tag=

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The invention describes a new antisense oligonucleotide, having a sequence comprising 8-80 bp targeted to a nucleic acid encoding cyclindependent kinase 4, specifically hybridises with the nucleic acid encoding cyclin-dependent kinase 4 and inhibits expression of cyclindependent kinase 4. The antisense oligonucleotide is useful for preparing a composition for treating diabetes, infertility or hyperproliferative disorder, e.g., cancer This sequence represents a human cyclin dependent
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           _base= OTHER
s= "OTHER= Phosphorothioate backbone. All cytidines
                                                                                                                                                                                                                                                                                                           New antisense oligonucleotide, having a sequence targeted to a nucleic acid encoding cyclin-dependent kinase 4, useful for preparing a composition for treating diabetes, infertility or hyperproliferative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /mod_base= OTHER
/note= "OTHER= 2'-O-Methoxyethyl (2'-MOE) nucleotides"
/note= "OTHER= 2'-O-Methoxyethyl (2'-MOE) nucleotides"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              nucleotides"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   gene therapy, antisense technology, PCTAIRE protein kinase 2, neurological disorder; human, PCTAIRE protein kinase 2; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human PCTAIRE protein kinase 2 antisense oligonucleotide #47.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (2'-MOE)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Seguence 20 BP; 5 A; 4 C; 7 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    الابادية - "CHER"
/mot.e= "OTHER= 2'-O-Methoxyethyl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 14.2; DB 1;
Pred. No. 8.7e+02;
0; Mismatches 3;
                                                                                                                                                                                                                                                                                                                                                                                                                  Example 15; SEQ ID NO 106; 90pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /note= "OTHER= Phospho:
are 5-methylcytidines"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              kinase 4 antisense oligonucleotide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      687 CAACCTTGTGGCACTCAAG 705
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           20 ccacrirérésécercaas 2
                                                                                                                                                                                                                                       Dobie KW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              BP.
                                                                                                                  02-JUL-2002; 2002US-00188779
                                                                                                                                                       02-JUL-2002; 2002US-00188779.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADI19193 standard; DNA; 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity 84.2'
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   15. .20
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/*tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /*tag=
                                                                                                                                                                                                                                                                                                                                                                               disorder, e.g., cancer.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /mod
                                                                                                                                                                                                                                       Freier SM,
                                                                                                                                                                                                                                                                          WPI; 2004-081710/08
                                         US2004005567-A1
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modified_base
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                                                                             08-JAN-2004.
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                                                                                                                                                                                                                                       Dean NM,
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                                                                                                                                                                                                                                                                                                                                                    The invention describes a compound 8-80 mucleobases in length targeted to a nucleic acid molecule encoding phosphodiesterase 4D. The compound specifically hybridises with the nucleic acid molecule encoding phosphodiesterase 4D. The acid molecule encoding phosphodiesterase 4D, or specifically hybridises with at least an 8-nucleobase portion of an active site on a nucleic acid molecule encoding phosphodiesterase 4D. The expression of phosphodiesterase 4D. The expression of phosphodiesterase 4D, and for treating diseases or conditions associated with expression of phosphodiesterase 4D, e.g. cancer, cardiovascular disease or inflammation. The antisense compounds are also useful as research reapplications, e.g. to prevent or delay infection, inflammation or tummour formation. This sequence represents a
                                                                                                                                                                                                               New antisense compounds targeted to nucleic acid molecules encoding phosphodiesterase 4D, useful for treating diseases associated with expression of phosphodiesterase 4D, e.g. cancer, cardiovascular disease
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /note= "OTHER= Phosphorothioate backbone. All cytidines are 5-methylcytidines"
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/note== "OTHER= 2'-O-Methoxyethyl (2'-MOE) nucleotides"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0.8%; Score 14.2; DB 1; Length 20;
84.2%; Pred. No. 8.7e+02;
iive 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            cytostatic; antidiabetic; antiinfertility; gene therapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Cyclin dependent kinase 4 antisense oligonucleotide #87.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   numan phosphodiesterase 4D antisense oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 20 BP; 5 A; 6 C; 7 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                       Example 15; SEQ ID NO 46; 72pp; English.
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                                                                                                                                           Roach MP;
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/mod base= OTHER
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                      15-MAY-2002; 2002US-00146860
                                                             15-MAY-2002; 2002US-00146860
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Best Local Similarity 84.2
Matches 16; Conservative
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                                                                                                                                           Dobie KW,
                                                                                                    (ISIS-) ISIS PHARM INC
                                                                                                                                                                              WPI; 2004-060214/06.
                                                                                                                                                                                                                                                                                     or inflammation
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modified_base
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                                                                                                                                           Bennett CF,
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 31-MAY-2002; 2002US-00160787.
                   31-MAY-2002; 2002US-00160787.
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                                                                                                                                                                                                                                                                 Local Similarity
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                                      (ISIS-) ISIS PHARM
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modified_base
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                                                                                                                                                                                            The invention describes a new antisense oligonucleotide, having a sequence comprising 8-80 bp targeted to a nucleic acid encoding PCTAIRE protein kinase 2, that specifically hybridises with the nucleic acid encoding PCTAIRE protein kinase 2 and having a sequence comprising 20 bp. The antisense oligonucleotide is useful for preparing a composition for treating e.g., neurological disorders. This sequence represents a human PCTAIRE protein kinase 2 antisense oligonucleotide.
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/note= "OTHER= Phosphorothioate backbone. All cytidines
are 5-methylcytidines"
                                                                                                                                      nucleic
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/mod_base= OTHER
/note= "OTHER= 2'-O-Methoxyethyl (2'-MOE) nucleotides"
/sociation
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                                                                                                                                   New antisense oligonucleotide, having a sequence targeted to a acid encoding PCTAIRE protein kinase 2, useful for preparing a composition for treating neurological disorders.
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84.2%; Pred. No. 8.7e+02;
tive 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                         Sequence 20 BP; 4 A; 1 C; 8 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                        Claim 1; SEQ ID NO 60; 58pp; English
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                                     31-MAY-2002; 2002US-00160787.
                                                        31-MAY-2002; 2002US-00160787
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                                                                           (ISIS-) ISIS PHARM INC
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les 16; Conserv
US2003225256-A1
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                                                                                                                                                                                                                                                                                            Query Match
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Matches
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New antisense oligonucleotide, having a sequence targeted to a nucleic acid encoding PCTAIRE protein kinase 2, useful for preparing a composition for treating neurological disorders.
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                                                                                                                                                Claim 1; SEQ ID NO 68; 58pp; English
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/note= "OTHER= P
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/mod_base= OTHER
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/mod_base= OTHER
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The invention describes a new antisense oligonucleotide, having a sequence comprising 8-80 bp targeted to a nucleic acid encoding PCTAIRE protein kinase 2, that specifically hybridises with the nucleic acid encoding PCTAIRE protein kinase 2 and having a sequence comprising 20 bp. The antisense oligonucleotide is useful for preparing a composition for treating e.g., neurological disorders. This sequence represents a human PCTAIRE protein kinase 2 antisense oligonucleotide.
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/note= "OTHER= Phosphorothioate backbone. All cytidines
are 5-methylcytidines"
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                                 nucleic
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                               New antisense oligonucleotide, having a sequence targeted to a acid encoding PCTAIRE protein kinase 2, useful for preparing a composition for treating neurological disorders.
                                                                                                                                                                                                                                                                                                                                      0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human PCTAIRE protein kinase 2 antisense oligonucleotide #52.
                                                                                                                                                                                                                                                                                                   Length 20;
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                                                                                                                                                                                                                                                                     Sequence 20 BP; 4 A; 3 C; 4 G; 9 T; 0 U; 0 Other;
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/mod_base= OTHER
/note= "OTHER= 2'-0-Methoxyethyl
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/note= "OTHER= 2'-O-Methoxyethyl
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Pred. No. 8.7e+02;
                                                                                                                                                                                                                                                                                                                                    0; Mismatches
                                                                                                   Example 15; SEQ ID NO 126; 58pp; English
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hes 16; Conservative
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WPI; 2004-022085/02
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neurological d
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                                                                                                  New antisense oligonucleotide, having a sequence targeted to a nucleic acid encoding PCTAIRE protein kinase 2, useful for preparing a composition for treating neurological disorders.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human PCTAIRE protein kinase 2 antisense oligonucleotide #113
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              antisense technology; PCTAIRE protein kinase 2; disorder; human; PCTAIRE protein kinase 2; ss.
                                                                                                                                                                                                                                                                                                                                                                       0.8%; Score 14.2; DB 1; Length 20;
84.2%; Pred. No. 8.7e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (2'-MOE)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (2'-MOE)
                                                                                                                                                                                                                                                                                                                                                                                                       Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /mod_base= OTHER
/note= "OTHER= 2'-O-Methoxyethyl
                                                                                                                                                                                                                                                                                                                                        BP; 6 A; 6 C; 4 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   -O-Methoxyethyl
                                                                                                                                                                                                                                                                                                                                                                                                          0; Mismatches
                                                                                                                                                                    Example 15; SEQ ID NO 128; 58pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                          1256 TAGGAACCCCAACTGAGGA 1274
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1. .20
/*tag= b
/mod_base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2 TAGGAACTCCATCTCAGGA 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        31-MAY-2002; 2002US-00160787.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       31-MAY-2002; 2002US-00160787
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADI19259 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                entry)
                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PHARM INC
 (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first
                                                                WPI; 2004-022085/02
                                                                                                                                                                                                                                                                                                                                                                                       Local Similarity
Les 16; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      US2003225256-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              gene therapy;
neurological
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                modified base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   modified base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SISI (-SISI)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Homo sapiens
                                                                                                                                                                                                                                                                                                                                        Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              22-APR-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     04-DEC-2003
                                                                                                                                                                                                                                                                                                                                                                       Query Match
                                   Watt AT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Watt AT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1120
                                                                                                                                                                                                                                                                                                                                                                                                          Matches
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The invention describes a new antisense oligonucleotide, having a sequence comprising 8-80 bp targeted to a nucleic acid encoding PCTAIRE protein kinase 2, that specifically hybridises with the nucleic acid encoding PCTAIRE protein kinase 2 and having a sequence comprising 20 bp. The antisense oligonucleotide is useful for preparing a composition for treating e.g., neurological disorders. This sequence represents a human PCTAIRE protein kinase 2 antisense oligonucleotide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The present invention relates to an isolated human protein with cancersuppressing function (HOMO). The protein, its encoded polynucleotide and antagonist are applicable in diagnosis and treatment of various diseases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Novel human protein with cancer-suppressing function, encoded polynucleotide and antagonist, applicable in diagnosis and treatment of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              e.g. cancer, The present sequence represents a primer of the invention.
                                                                                                                                                                                   Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
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0
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Pred. No. 8.7e+02;
0; Mismatches 3; Indels
                                                                                                                                                       Length 20;
                                                                                                                                                                                 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                         cancer; Cytostatic; cancer; ds; HOMO; ss; primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 20 BP; 3 A; 5 C; 6 G; 6 T; 0 U; 0 Other;
                                                                                                                       Sequence 20 BP; 7 A; 8 C; 1 G; 4 T; 0 U; 0 Other;
                                                                                                                                                   Score 14.2; DB 1;
Pred. No. 8.7e+02;
0; Mismatches 3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 1; SEQ ID NO 73; 80pp; Chinese.
                                                                                                                                                                                                               988 CCCCAGAACCIGCTCAICA 1006
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   963 GAAGGTGCTACACCGAGAC 981
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                                                                                                                                                     0.8%;
                                                                                                                                                                                                                                              2 CCACAGAACCTCCTCATTA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             07-AUG-2003; 2003WO-CN000639.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        07-AUG-2002; 2002CN-00136400.
16-SEP-2002; 2002CN-00137000.
16-SEP-2002; 2002CN-00137009.
20-NOV-2002; 2002CN-00145435.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         various diseases e.g. cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GAAGGTGCTACTCCAAGCC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       84.2%;
                                                                                                                                                                                                                                                                                                                                                                                                             Primer of the invention #27
                                                                                                                                                                                                                                                                                                                     ADK00650 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Best_Local Similarity 84.2
Matches 16; Conservative
                                                                                                                                                                                   Conservative
                                                                                                                                                                                                                                                                                                                                                                                 (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2004-191733/18.
                                                                                                                                                                Local Similarity
nes 16; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (NEWO-) NEWORGEN
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 19-FEB-2004.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Synthetic
                                                                                                                                                     Query Match
                                                                                                                                                                                                                                                                                                                                                   ADK00650;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Yang S,
                                                                                                                                                                                                                                                                                        RESULT 1123
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                                                                                                                                                                                   Matches
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      88888888888
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                                                           The invention describes a new antisense oligonucleotide, having a sequence comprising 8-80 bp targeted to a nucleic acid encoding PCTAIRE protein kinase 2, that specifically hybridises with the nucleic acid encoding PCTAIRE protein kinase 2 and having a sequence comprising 20 bp. The antisense oligonucleotide is useful for preparing a composition for treating e.g., neurological disorders. This sequence represents a human PCTAIRE protein kinase 2 antisense oligonucleotide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /note= "OTHER= Phosphorothioate backbone. All cytidines
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New antisense oligonucleotide, having a sequence targeted to a nucleic acid encoding PCTAIRE protein kinase 2, useful for preparing a composition for treating neurological disorders.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         'note= "OTHER= 2'-0-Methoxyethyl (2'-MOE) nucleotides"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 nucleotides
                                                                                                                                                                                                                                              Gaps
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0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human PCTAIRE protein kinase 2 antisense oligonucleotide #109.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    kinase 2;
                                                                                                                                                                                                             0.8%; Score 14.2; DB 1; Length 20;
84.2%; Pred. No. 8.7e+02;
tive 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ; antisense technology; PCTAIRE protein kinase disorder; human; PCTAIRE protein kinase 2; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /mod_base= OTHER
/note= "OTHER= 2'-O-Methoxyethyl (2'-MOE)
                                                                                                                                                                                 Sequence 20 BP; 9 A; 4 C; 3 G; 4 T; 0 U; 0 Other;
      composition for treating neurological disorders.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example 15; SEQ ID NO 122; 58pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      are 5-methylcytidines
                                Claim 1; SEQ ID NO 65; 58pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Location/Qualifiers
                                                                                                                                                                                                                                                                        1172 GCATCTTCTATGAGATGGC 1190
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /mod_base= OTHER
                                                                                                                                                                                                                                                                                                    N
                                                                                                                                                                                                                                                                                           GCATTTTCTTTGAAATGGC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 31-MAY-2002; 2002US-00160787
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                                                                                                                                                                                                                                           Conservative
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                                                                                                                                                                                                                                                                                                                                                                               DNA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         mod/
                                                                                                                                                                                               Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                             ADI19255 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US2003225256-A1
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modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  therapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    modified base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               modified base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  neurological
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                                                                                                                                                                                                                                                                                                                                                   RESULT 1122
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     gene
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Hinrikson HP;

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Use of internal transcribed spacer 1 nucleic acid sequences, for distinguishing species of Aspergillus from one another, or detecting the presence of an Aspergillus species or strain of a species in a biological
                                                                                  (USSH ) US DEPT HEALTH & HUMAN SERVICES.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               16-MAY-2003; 2003WO-US016076.
                                  16-MAY-2003; 2003WO-US016076
                                                           17-MAY-2002; 2002US-0381463P
                                                                                                                                  WPI; 2004-022871/02.
                                                                                                            Morrison CJ,
          27-NOV-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADK23613;
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                                                                                                                                                                                                                                                                                                                                                                                                                    The invention relates to a novel method for detecting a specific white rot fungi strain from white rot fungi present in soil. The method comprises: obtaining a soil sample; recovering DNA from the soil using an elution buffer; purifying the DNA; preparing the DNA sample; determining strain specific base sequence in the genome DNA of the white rot fungi strain; preparing a primer; performing amplification using the prepared primer and the DNA sample as a template, and detecting the amplified white rot fungi strain DNA. The method enables detection of the specific white rot fungi strain from the white rot fungi strain from the white rot fungi strain from the white rot fungi present in the soil. This polymucleotide sequence represents a primer used in the exemplification
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Aspergillus; internal transcribed spacer; ITS; ITS1-V1; ITS-V2; ITS-V3;
                                                                                                                                                                                                                                                                                                                                   Detecting specific white rot fungi strain, comprises obtaining soil, recovering DNA using elution buffer, preparing DNA sample and primer, performing amplification using DNA sample as template and detecting
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         .
0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0.8%; Score 14.2; DB 1; Length 20; 84.2%; Pred. No. 8.7e+02; ive 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 20 BP; 2 A; 6 C; 5 G; 7 T; 0 U; 0 Other;
                                                                                   White rot fungi strain related primer, ITS-2R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Fungal universal forward primer #SEQ ID 29.
                                                                                                            white rot fungi; soil; ss; primer; ITS-2R
                                                                                                                                                                                                                                                           KAGA-) KAGAKU GIJUTSU SHINKO JIGYODAN.
                                                                                                                                                                                                                                                                                                                                                                                                 Claim 7; SEQ ID NO 3; 15pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1549 CTTCGGTCTTCGTCGATGC 1567
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ITS-V4; ITS-V5; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        crecerrerreares 20
            BP.
                                                                                                                                                                                                           09-JUL-2002; 2002JP-00200262
                                                                                                                                                                                                                                     09-JUL-2002; 2002JP-00200262
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            ADL16005 standard; DNA; 20
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                                                            (first entry)
                                                                                                                                                                                                                                                                        (KOND/) KONDO R.
(WAKA-) WAKAO KENSETSU KK.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match 0.8
Best Local Similarity 84.2
Matches 16; Conservative
                                                                                                                                                                                                                                                                                                            WPI; 2004-242914/23.
                                                                                                                                                                                                                                                                                                                                                                          amplified sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  the invention.
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                                                                                                                                                             JP2004041024-A.
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                                                            06-MAY-2004
                                                                                                                                                                                    12-FEB-2004
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                                     ADL16005;
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 ADL16005
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The invention relates to a method for distinguishing species of
Aspergillus from one another, comprising detecting differences in two or
more internal transcribed spacer (ITS), i.e. ITS1-V1, ITS-V2, ITS-V3, ITS
-V4 and ITS-V5, nucleic acid sequences of Aspergillus. The internal
transcribed spacer 1 nucleic acid sequences are useful for distinguishing
species of Aspergillus from one another, or for detecting the presence of
species of Aspergillus from an another, or for detecting the presence of
invention, a biological sample was obtained from an infected subject, and
invention, a biological sample was obtained from an infected subject, and
invention, a biological sample was obtained from an infected subject, and
isolation of fungal DNA. The universal pairs internal transcribed spacer
(ITS)s and ITS2, ITSS and ITS4, ITS1 and ITS2, or ITS1 and ITS4 were
added to the reaction mixture to amplify the fungal DNA present in the
mixture. PCR products generated with the primer pairs were purified, and
the purified products were sequenced on both strands using the same
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ó
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                primers as initially used for PCR amplification. Sequencing products were purified and analysed on an automated capillary DNA sequencer. A total of 46 ITS1 consensus sequences ranging in length from 142-187 nucleotides were compiled. Overall, 5 regions with significant interspecies variability in length and sequence were recognised. The current sequence represents the nucleic acid sequence of a fungal universal forward
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Aspergillus, internal transcribed spacer; ITS; ITS1-V1; ITS-V2; ITS-V3; ITS-V4; ITS-V5; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 20 BP; 2 A; 6 C; 5 G; 7 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0; Mismatches
Disclosure; SEQ ID NO 29; 56pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1549 CTTCGGTCTTCGTCGATGC 1567
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The invention relates to a method for distinguishing species of
Aspergillus from one another, comprising detecting differences in two or
more internal transcribed spacer (ITS), i.e. ITS1-V1, ITS-V2, ITS-V3, ITS-V3, ITS-V3, TS-V3, TS-
                                                                                                                                                                                                                                                                                                                        Use of internal transcribed spacer 1 nucleic acid sequences, for distinguishing species of Aspergillus from one another, or detecting the presence of an Aspergillus species or strain of a species in a biological
                                                                                                     (USSH ) US DEPT HEALTH & HUMAN SERVICES.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Disclosure; SEQ ID NO 32; 56pp; English.
                                     17-MAY-2002; 2002US-0381463P.
                                                                                                                                                                                 Morrison CJ, Hinrikson HP;
                                                                                                                                                                                                                                                    WPI; 2004-022871/02.
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0.8%; Score 14.2; DB 1; Length 20; 84.2%; Pred. No. 8.7e+02; 3; Indels Sequence 20 BP; 7 A; 5 C; 6 G; 2 T; 0 U; 0 Other; 0; Mismatches 1549 CTTCGGTCTTCGTCGATGC 1567 Query Match
Best Local Similarity 84.2*
Matches 16; Conservative

crecerretrearcearee 1 13 g

ADL06765 standard; DNA; 20 BP 06-MAY-2004 (first entry) ADL06765; ADLO6765/ XXX ADLO0 XXX ADLO0 XXX ADLO0 XXX Human XXX Human XXX Human XXX Human XXX Human XXX Homo XXX Homo

RESULT 1127

Factor VII variant allele-specific probe/PCR primer F710G, SEQ 36

Human; factor VII; FVII; chromosome 11q34-qter; allelic variation; blasma stability; function; single nucleotide polymorphism; SNP; insertion; predisposition; cardiovascular disease; detection; diagnosis; patient-specific treatment; coagulation disorder; thrombotic disorder; thrombosis; haemostatic; thrombolytic; allele-specific; PCR; primer; probe; ss.

Homo sapiens.

WO2004011675-A2.

05-FEB-2004

23-JUL-2003; 2003WO-ES000379.

for factor VII-related disorders. Additionally, factor VII proteins containing at least one allelic variation of the invention (e.g., one which increases plasma stability) are useful as pharmaceuticals for treating disorders of blood coagulation or thrombosis. The present sequence represents a specifically claimed allelic variant-specific oligonuclectide probe, which was also used as a PCR primer in an example New allelic variants of the factor VII gene, useful in diagnosing predisposition to cardiovascular disease and for treatment of e.g. coagulation disorders and thrombosis. The invention relates to a human factor VII (FVII) polynucleotide (PRIV-) FUNDACIO PRIVADA I INST RECERCA LHOSPITA. Soria Fernandez JM, Claim 7; SEQ ID NO 36; 32pp; Spanish. 25-JUL-2002; 2002ES-00001749. WPI; 2004-143875/14 Fontcuberta Boj J, of the invention.

Gaps ; 0 0.8%; Score 14.2; DB 1; Length 20; 84.2%; Pred. No. 8.7e+02; ive 0; Mismatches 3; Indels Sequence 20 BP; 3 A; 5 C; 5 G; 7 T; 0 U; 0 Other; 16; Conservative Query Match Best Local Similarity Matches

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Gaps

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RESULT 1128

ADJ45621 standard; DNA; 20 BP 06-MAY-2004 (first entry) ADJ45621; ADJ45621/

Human GPCR 12 antisense oligonucleotide ISIS238089.

Human, 88, antisense gene therapy, G protein-coupled receptor 12, GPCR 12, appetite control, nootropic, neuroprotective, neuroendocrine system disorder, signal transduction disorder; neuronal disorder; motor disorder; sensory disorder; psychiatric disorder; behavioural disorder.

Homo sapiens.

/mod_base= OTHER /note= "All cytidines are 5-methylcytidines and all Location/Qualifiers Д 1. .20 /*tag= modified base

stability and/or function of the gene or of the encoded factor VII. 49 stability and/or function of the gene or of the encoded factor VII. 49 catorial or via a stability and/or function of the gene or of the encoded factor VII. 49 cator VII allele with a so include insertions, are tabulated in the specification and are indicative of a predisposition to cardiovascular disease. Factor VII activates the coagulation cascade after it binds to tissue factor, via the encoding it is located on chromosome 13q34-qter. The invention also relates to 36 oligonucleotide or probes specific for the variant factor VII alleles (AbL06730-AbL06765), a process for testing for the presence of any of the 49 variations, and a diagnostic kit which contains the allele-specific oligonucleotides. The method of the invention can be used to detect the presence of variant cator VII alleles, and thus to determine whether an individual has a predisposition to cardiovascular disorders. Analysis of the factor VII alleles variations will permit the design of patient-specific treatments

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/note= "This oligonucleotide has a phosphorothioate backbone and 2-'methyoxyethyl (2'-MOE) wings at the 5' and 3' ends, which are 4 nucleotides in length. Also all
                                                                                                                          Antilipaemic, Cardiovascular, Analgesic, Antianginal, Antisense therapy, Human, Endothelial Lipase, dyslipidaemia, high density lipoprotein, HDL, cardiovascular disorder, metabolic syndrome X; ss.
                                                                           Human endothelial lipase antisense oligonucleotide, SEQ ID 1053.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              cytidine residues are 5-methylcytidines"
                                                                                                                                                                                                                                                                                                                        Location/Qualifiers
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/mod_base= OTHER
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                      20-MAY-2004 (first entry)
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                                                                                                                                                                                                                                                                                                                                                  modified base
                                                                                                                                                                                                                                             Homo sapiens
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                                                                                                                                                                                                                                                                      Synthetic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The invention relates to a compound 8-80 nucleobases in length targeted to, and which specifically hybridises with a nucleic acid molecule encoding GPCR-12 (protein-coupled receptor 12), and inhibits the expression of GPCR-12, i.e. an antisense oligonucleotide (AS). Also are a compound 8-80 nucleobases in length that specifically hybridises with at least an 8-nucleobase portion of an active site on a nucleic acid molecule encoding GPCR-12, a composition comprising the compound and a carrier or diluent, inhibiting the expression of GPCR-12 in cells or tissues with the compound so that expression of GPCR-12 is inhibited), treating an animal having a disease or condition associated with GPCR-12 (by administering to the animal at the rapeutic or prophylactic amount of the compound (by contacting a preferred target region of a nucleic acid molecule encoding GPCR-12 with one or more candidate antisense compounds that inhibit or and selecting for one or more candidate antisense compounds that inhibit the expression of a nucleic acid encoding GPCR-12). The compound, composition and methods are useful for treating a disease or condition and methods are useful for treating a disease or condition in present compounds that inhibit composition and methods are useful for treating a disease or condition and methods are useful for treating a disease or condition and methods are useful for treating a disease or condition in tissue, and selecting the system of a disease or condition in breat tissue, and selecting the system or an advance of the preferred target region and methods are useful for treating a disease or condition in tissue, and selecting the system of the preferred target and preferred target and part of the preferred target
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New compounds, particularly antisense oligonucleotides targeted to a nucleic acid encoding G protein-coupled receptor 12 (GPCR-12), useful for treating diseases of the neuroendocrine system.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
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inkages are phosphorothioate linkages"
                                                                                                                                                                                  /mod_base= OTHER
/note= "2'-methoxyethyl-nucleotide'
                                                                                                       'note= "2'-methoxyethyl-nucleotide'
16. .20
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Dobie KW;
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                           modified_base
                                                                                                                                   modified_base
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New antisense oligonuclectide for modulating endothelial lipase expression, for diagnosing, preventing or treating e.g. dyslipidemia, low high density lipoprotein or cardiovascular disorders.
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                                                                                                                                                                                 The present invention relates to antisense oligonucleotides (AbJ21603-AbJ25510) targeted to human Endothelial Lipase (EL) coding sequence (AbJ25517), where the antisense oligonucleotide specifically hybridises with and inhibits the expression of EL. The antisense oligonucleotides are useful for modulating the expression of endothelial lipase in cells or tissues to treat diseases associated with EL expression, such as dyslipidaemia, low high density lipoprotein (HDL), cardiovascular disorder or metabolic syndrome X. In addition, the oligonucleotides are
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                         used for diagnostics, prophylaxis, or as research reagents or kits.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human endothelial lipase antisense oligonucleotide, SEQ ID 803.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 14.2; DB 1; Length 20; Pred. No. 8.7e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 20 BP; 5 A; 2 C; 10 G; 3 T; 0 U; 0 Other;
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                                                                                                                               Claim 3; SEQ ID NO 1053; 1007pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1689 CITCCCIGCTIACTCTCTG 1707
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       20 CTCCCCAGCTCACTCTCTG 2
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Best Local Similarity 84.2%;
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ADJ22655 standard; DNA; 20 BP.

RESULT 1129 ADJ22655 AXH AXH

В ð

ADJ22655

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expression, for diagnosing, preventing or treating e.g. dyslipidemia, low high density lipoprotein or cardiovascular disorders.
                                                                                    /note= "This oligonucleotide has a phosphorothioate backbone and 2-'methyoxyethyl (2'-MOE) wings at the 5' and 3' ends, which are 4 nucleotides in length. Also all
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The present invention relates to antisense oligonucleotides (ADJ21603-ADJ25510) targeted to human Endothelial Lipase (EL) coding sequence (ADJ25517), where the antisense oligonucleotide specifically hybridises with and inhibits the expression of EL. The antisense oligonucleotides are useful for modulating the expression of endothelial lipase in cells or tissues to treat diseases associated with EL expression, such as dyslipidaemia, low high density lipoprotein (HDL), cardiovascular disorder or metabolic syndrome X. In addition, the oligonucleotides are used for diagnostics, prophylaxis, or as research reagents or kits.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New antisense oligonucleotide for modulating endothelial lipase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0.8%; Score 14.2; DB 1; Length 20;
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0; Mismatches 3; Indels
                                                                                                                                                       cytidine residues are 5-methylcytidines
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    Location/Qualifiers
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                                                                /mod_base= OTHER
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                                                                                                                                                                                                                                                                                                                                 19-JUL-2002; 2002US-0397106P
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nes 16; Conservative
                                             *tag=
                                                                                                                                                                                                                                                                                                                                                                            (PHAA ) PHARMACIA CORP
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    Key
modified base
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modified_base
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                                                                                                                                                                                                                                                                                                                                                                                                                          Bhat BG;
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    δ
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New antisense oligonucleotide for modulating endothelial lipase expression, for diagnosing, preventing or treating e.g. dyslipidemia, low high density lipoprotein or cardiovascular disorders.
                                                                                                                                                                                                                   /note= "This oligonucleotide has a phosphorothioate backbone and 2-'methyoxyethyl (2'-MOE) wings at the 5' and 3' ends, which are 4 nucleotides in length. Also all
Human; Endothelial Lipase; dyslipidaemia; high density lipoprotein; HDL; cardiovascular disorder; metabolic syndrome X; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Antilipaemic; Cardiovascular; Analgesic; Antianginal; Antisense therapy; Human; Endothelial Lipase; dyslipidaemia; high density lipoprotein; HDL; cardiovascular disorder; metabolic syndrome X; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The present invention relates to antisense oligonucleotides (AbJ21603-AbJ25510) targeted to human Endothelial Lipase (EL) coding sequence (AbJ25517), where the antisense oligonucleotide specifically hybridises with and inhibits the expression of EL. The antisense oligonucleotides are useful for modulating the expression of endothelial lipase in cells or tissues to treat diseases associated with EL expression, such as dyslipidaemia, low high density lipoprotein (HDL), cardiovascular disorder or metabolic syndrome X. In addition, the oligonucleotides are used for diagnostics, prophylaxis, or as research reagents or kits.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human endothelial lipase antisense oligonucleotide, SEQ ID 3650
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0
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84.2%; Pred. No. 8.7e+02;
ative 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                     cytidine residues are 5-methylcytidines"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 3; SEQ ID NO 803; 1007pp; English.
                                                                                                                                location/Qualifiers
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                                                                                                                                                                                                   /mod base= OTHER
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nes 16; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (PHAA ) PHARMACIA CORP
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                                                                                                                                Key
modified base
                                                                sapiens
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                                                                                       Synthetic
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BG; Bhat

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Gaps

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/note= "This oligonucleotide has a phosphorothioate backbone and 2-'methyoxyethyl (2'-MOE) wings at the

/mod_base= OTHER

Homo sapiens

19

à g

Query Match

ADJ25252;

ADJ25252,

Synthetic

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nucleic acid molecule encoding Navi.3, where the antisense compound specifically hybridizes with and inhibits the expression of Navi.3. The compound and composition are useful for treating a disease or condition associated with Navi.3, e.g. pain including but not limited to neuropathic pain, post-herpetic neuralgia, chronic pain, lower back pain, diabetic neuropathy, trigeminal neuropathy arthritic pain, lower back pain, abain from burns, migraine headache, cluster headache, mild-to-moderate headache; seizure disorder such as childhood seizure disorder; including but not limited to neonatal or infantile epilepsy; or ataxia. The present sequence represents a chimeric phosphorothhoate oligomucleotide with 2'MOE wings and a deoxy gap. Used during the antisense inhibition of human Navi.3 expression, the oligomucleotides are designed to target different regions of the human Navi.3 RNA.
                                  Navl.3, useful for useful for treating a disease or condition associated with Navl.3, e.g. pain, seizure disorder such as childhood seizure disorder, or ataxia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New antisense compound targeted to a nucleic acid molecule encoding Navl.3, useful for useful for treating a disease or condition associated with Navl.3, e.g. pain, seizure disorder such as childhood seizure disorder, or ataxia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Navl.3; Analgesic; Nootropic; Neuroprotective; post-herpetic neuralgia; diabetic neuropathy; arthritic pain; migraine headache; infantile epilepsy; ataxia; ss.
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                   New antisense compound targeted to a nucleic acid molecule encoding
                                                                                                                                                                                     present invention relates to an antisense compound targeted to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Chimeric phosphorothioate oligonucleotide to target Navl.3 #1553
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0.8%; Score 14.2; DB 1; Length 20;
14.2%; Pred. No. 8.7e+02;
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                                                                                                                                         Claim 4; SEQ ID NO 1275; 417pp; English.
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Best Local Similarity
Matches 16; Conserv
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ADK74219/c
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                                                                                                                                                                                                                                                                                                                                                                                                     New antisense oligonucleotide for modulating endothelial lipase expression, for diagnosing, preventing or treating e.g. dyslipidemia, low high density lipoprotein or cardiovascular disorders.
and 3' ends, which are 4 nucleotides in length. Also all cytidine residues are 5-methylcytidines"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADJ25510) targeted to human Endothelial Lipase (EL) coding sequence (ADJ25517), where the antisense oligonucleotide specifically hybridises with and inhibits the expression of EL. The antisense oligonucleotides are useful for modulating the expression of endothelial lipase in cells dyslipidaemia, low high density lipopototic (HDL), cardiovascular disorder or metabolic syndrome X. In addition, the oligonucleotides are used for diagnostics, prophylaxis, or as research reagents or kits.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The present invention relates to antisense oligonucleotides (ADJ21603-
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Chimeric phosphorothioate oligonucleotide to target Nav1.3 #1275.
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84.2%; Pred. No. 8.7e+02;
tive 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 20 BP; 1 A; 6 C; 6 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 3; SEQ ID NO 3853; 1007pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1375 GACGGGCCGACCTCCTCA 1393
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     infantile epilepsy; ataxia; ss.
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Best Local Similarity 84.2
Matches 16, Conservative
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specifically hybridizes with and inhibits the expression of Navl.3. The compound and composition are useful for treating a disease or condition associated with Navl.3. e.g. pain including but not limited to neuropathic pain, post-herpetic neuralgia, chronic pain, lower back pain, diabetic neuropathy, trigeminal neuropathy, arthritic pain, acute pain, pain from burns, migraine headache, cluster headache, mild-to-moderate headache; seizure disorder such as childhood seizure disorder such as childhood seizure disorder such se childhood seizure disorder including sequence represents a chimeric phosphorothioate oligomucleotide with 2.MOE wings and a deoxy gap. Used during the antisense inhibition of human Navl.3 expression, the oligomucleotides are designed to target
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The invention relates to an antisense compound targeted to a nucleic acid molecule encoding human vascular endothelial growth factor (VECF) corregulated chemokine-1 (VCC-1), and which specifically hybridises with and inhibits the expression of VCC-1. The invention also relates to a composition comprising the antisense compound, a method of inhibiting the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 immunological disorder; cardiovascular disorder; neurological disorder; ischaemia; reperfusion injury; cancer; angiogenic disorder; haemangioma; tumour angiogenesis; rheumatoid arthritis; atherosclerosis; psoriasis; fibrosis; myocardial infarction; wound healing; bone fracture;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human VEGF co-regulated chemokine-1 DNA antisense oligonucleotide #441.
                                                                                                                                                                                                                                                                                                                                                                       Gaps
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                                                                                                                                                                                                                                                                                                                           0.8%; Score 14.2; DB 1; Length 20;
84.2%; Pred. No. 8.7e+02;
tive 0; Mismatches 3; Indels
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wscular endochelial growth factor; ss; antisense compound;
phosphorothioate linkage; 2'-O-methoxyethyl sugar moiety;
5-methylcytosine; antisense oligonucleotide; diabetes;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         cartilage damage; tissue regeneration; organ regeneration; periodontal disease; gut regeneration; atrial fibrillation.
                                                                                                                                                                                                                                                                                   Sequence 20 BP; 8 A; 1 C; 6 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                   human Navī.3 expression, the oligonucleotio
different regions of the human Nav1.3 RNA.
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nes 16; Conservative
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expression of VCC-1 in cells or tissues comprising contacting the cells or tissues with the antisense compound and a method of treating a human having a disease or condition associated with VCC-1 comprising administering the antisense compound to an animal to inhibit expression of VCC-1. The antisense oligomuclectide comprises at least one modified internucleoside linkage, preferably a phosphorothioate linkage. It also comprises at least one modified sugar moiety, preferably a 2'-0-comprises at least one modified sugar moiety, preferably a 2'-0-comprises at least one modified sugar moiety, and at least one modified nucleobase, specifically a 5-methylcytosine. The antisense compound is useful for it a chimeric oligomucleotide. The antisense compound is useful for treating a disease or condition associated with VCC-1, such as diabetes, an immunological disorder, a cardiovascular disorder, a neurological disorder, a cardiovascular disorder, an enuclogical disorder, a cardiovascular disorder or an anglogenic disorder, ischaemia, reperfusion injury, cancer or an anglogenic atherosclerosis, psociasis or tibrosis after myocardial infarction. VCC-1 antisense oligomucleotides may also be used for wound healing, for regeneration of the part of the contraction of contraction of contracting of contraction of contraction of contraction of contraction of
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 tissues or organs, for treating periodontal diseases, for gut protection or regeneration, for treatment of lung or liver fibrosis or for management of atrial fibrillation. This sequence represents an antisense
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human VEGF co-regulated chemokine-1 DNA antisense oligonucleotide #482.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          oligonucleotide targeted to DNA encoding the human VCC-1 polypeptide
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match 0.8%; Score 14.2; DB 1; Length 2 Best Local Similarity 84.2%; Pred. No. 8.7e+02; Matches 16; Conservative 0; Mismatches 3; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             the invention.
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Tchernev VT, Spytek KA, Zerhusen BD, Patturajan M, Shimkets RA;
Li L, Gangolli EA, Padigaru M, Anderson DW, Rastelli L, Miller CE;
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2001US-0288527P.
2001US-0288504P.
2001US-0294473P.
2001US-0296564P.
2001US-0298959P.
2001US-0298959P.
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2001US-0312908P
2001US-031390P
2001US-0315470P
2001US-0316447P
2001US-0318115P
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2001US-0278775P.
2001US-0278778P.
2001US-0279882P.
2001US-0279884P.
2001US-0280147P.
2001US-0282992P.
2001US-028393P.
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2001US-0323379P.
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2001US-0330308P.
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GANGOLLI E A.
PADIGARU M.
ANDERSON D W.
RASTELLI L.
MILLER C E.
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ZERHUSEN B D.
PATTURAJAN M.
SHIMKETS R A.
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TAUPIER R J.
GUSEV V Y.
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WOLENC R R.
PENA C E A.
FURTAK K.
GROSSE W M.
ALSOBROOK J E
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26-FEB-2001; 2
27-FEB-2001; 2
27-FEB-2001; 2
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14-MAR-2001; 2
14-MAR-2001; 2
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08-JUN-2001;
18-JUN-2001;
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15-MAR-2001;
16-MAR-2001;
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29-MAY-2001;
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02-MAR-2001;
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28-AUG-2001;
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07-SEP-2001;
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18-OCT-2001;
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(RIEG/)
(BURG/)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Li L,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GUSE/
The invention relates to an antisense compound targeted to a nucleic acid molecule encoding human vascular endothelial growth factor (VEGF) coregulated chemokine-1 (VCC-1), and which specifically hybridises with and inhibits the expression of VCC-1. The invention also relates to a composition comprising the antisense compound, a method of inhibiting the expression of VCC-1 in cells or tissues compound and a method of treating the cells or tissues with the antisense compound and a mathod of treating a human core expression of VCC-1. The antisense compound to an animal to inhibit expression of volving a disease or condition associated with VCC-1 comprising a human core of VCC-1. The antisense oligoncolectide comprises at least one modified sugar moiety, preferably a 2'-0-comprises at least one modified sugar moiety, preferably a 2'-0-comprises at least one modified sugar moiety, preferably a 2'-0-comprises at least one modified sugar moiety, preferably a 2'-0-comprises at least one modified moiety, and at least one modified molectide. The antisense oligonucleotide. The antisense oligonucleotide of comprises of comprises of comprises of condition associated with VCC-1, such as diabetes, an immunological disorder, a cardiovascular disorder, a neurological disorder, a cardiovascular disorder, a neurological disorder, a cardiovascular disorder, an englogenic disorder, e.g. haemangioma, tumour angiogenesis, rhemmatoid arthritis, continense oligonucleotides may also be used for wound healing, for treatment of this sequence represents an antisense consense oligonucleotides may also be used for wound healing, for treatment of lung or liver fibrosis or for management of atrial fibrillation. This sequence represents an antisense oligonucleotide targeted to DNA encoding the human VCC-1 polypoptide of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ó
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human; ss; NOVX; cancer; diabetes; cardiomyopathy; atherosclerosis; PCR;
primer; RTQ PCR; real time quantitative PCR.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
 useful for treating VCC-1-associated disorders, e.g. diabetes or
neurologic disorder.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0;
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Pred. No. 8.7e+02;
0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Seguence 20 BP; 4 A; 7 C; 4 G; 5 T; 0 U; 0 Other;
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                                         Claim 4; SEQ ID NO 482; 336pp; English
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2001US-0265412P.
2001US-0265514P.
2001US-026517P.
2001US-0265406P.
2001US-026640FP.
2001US-026695P.
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Best Local Similarity 84.2%;
Matches 16; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                               the invention.
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07-FEB-2001;
07-FEB-2001;
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31-JAN-2001;
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02-FEB-2001;
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Homo sapiens
                                                                               17-JUN-2004
 Furtak K,
                                                                                            Synthetic.
                                                                           ADN06167;
                                                        Query Match
                                                                     RESULT 1138
                                                         Best Loca
Matches
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05-SEP-2003; 2003US-00655847. 31-MAY-2002; 2002US-00160807 JS2004063129-A1. modified base 01-APR-2004. 01-JAN-2004 invention. or tumors Synthetic ADL34826; Query Match RESULT 1139 Matches ADL34826 à ద ö Including splice variants) and the nucleic acids (NA) that encode them. Also included are the mature NOVX proteins (and their encoding polynucleotides), a vector comprising NOVX NA, a call comprising the vector, an antibody that binds immunospecifically to NOVX, determining the presence or amount of NOVX in a sample, determining the presence or amount of NOVX in a sample, determining the presence or modulating the activity of NOVX, treating or preventing a NOVX-associated disorder, determining the presence of or predisposition to a disease associated with altered levels of NOVX and treating a pathological state in a mammal comprising administering a polypeptide which is at least 95% identical to NOVX (or fragment). NOVX and NA may be used in the prevention, treatment and diagnosis of disease associated with Wolenc AR, Pena CEA; , Rieger DK, Burgess CE; inappropriate expression and activity of NOVX (e.g. cancer, diabetes, cardiomyopathy and/or atherosclerosis). The anti-NOVX antibodies and antagonists may also be used to down regulate expression and activity of NOVX. The anti-NOVX antibodies may also be used as diagnostic agents for detecting the presence of NOVX in samples (e.g. by enzyme linked immunosorbant assay (ELISA). The agents and methods may be used in this way to prevent, diagnose and treat cancer, diabetes, cardiomyopathy and/or atherosclerosis. The present sequence is a real time quantitative PCR (RTQ PCR) primer for tissue specific expression studies for a NOVX /mod_base= OTHER /note= "Phosphorothioate backbone in which all cytidines Isolated NOVX polypeptides and polymucleotides, useful for preventing diagnosing and/or treating cancer, diabetes, cardiomyopathy and atherosclerosis. Gaps Selenophosphate synthetase 2; SPS2; rheumatoid arthritis; infection; inflammation; tumour; antisense therapy; human; antisense; phosphorothioate backbone; ss. invention relates isolated 162 NOVX polypeptides (NOV1-NOV99 ; 0 0.8%; Score 14.2; DB 1; Length 20; ilarity 84.2%; Pred. No. 8.7e+02; Conservative 0; Mismatches 3; Indels Human SPS2 specific antisense oligonucleotide, ISIS 138236 /mod_base= OTHER /note= "2'-methoxyethyl nucleotides" Alsobrook JP, Lepley DM, Sequence 20 BP; 6 A; 5 C; 6 G; 3 T; 0 U; 0 Other; Colman SD, Example 2; SEQ ID NO 1149; 1309pp; English. are 5-methylcytidines" 1. .5 Location/Qualifiers Gusev VY, 506 AGGCTACCTGGAGAAGCT 524 20 BP 2 AGGACCATCTGGAGAGCT ADN06167 standard; DNA; 20 (first entry) Ω Taupier RJ, /*tag= /*tag= Grosse WM, WPI; 2004-180039/17. Local Similarity es 16; Conserv Key modified_base modified base Gerlach V,

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composition comprises antisense oligomodeotides targeted to SPS2 gene. The antisense oligomodeotide is useful for modulating the expression of SPS2 in cells or tissues to treat diseases associated with their expression, e.g. rheumatoid arthritis, infections, inflammation or tumours. It is also used for diagnostics, prophylaxis, or as research therappy. The present sequence is an antisense oligomodeotide targeted to human SPS2 DNA. This sequence is used in the exemplification of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The invention relates to antisense compounds, compositions and methods
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New antisense oligonucleotides for modulating selenophosphate synthetase 2 (SPS2) expression, useful for diagnosing, preventing or treating conditions associated with SPS2, e.g. rheumatoid arthritis, inflammation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
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phosphorothioate linkage; 2'-O-methoxyethyl sugar; 5-methylcytosine;
hyperproliferative disorder; cancer; cytostatic; gene therapy; ss;
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                                                                                                /note= "2'-methoxyethyl nucleotides"
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RESULT 1141
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                                                                                                                                                      This invention describes novel antisense oligonucleotides targeted to a nucleic acid encoding PPAR-delta, which specifically hybridise to and inhibit expression of PPAR-delta, The oligonucleotide specifically hybridises with at least an 8-nucleobase portion of an active site on the nucleic acid molecule encoding the PPAR-delta and comprises at least one modified internucleoside linkage, which is a phosphorothicate linkage, at least one modified sugar moiety, which is a 2-0-methoxyethyl sugar. The antisense oligonucleotides are useful for preparing a composition for treathing hyperproliferative disorders, e.g., cancer. The oligonucleotides of the invention have cytostatic activity and can be used for gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New antisense oligonucleotide, having a sequence targeted to a nucleic acid encoding PPAR-delta, useful for preparing a composition for treating
                                                                                             New antisense oligonucleotide, having a sequence targeted to a nucleic acid encoding PPAR-delta, useful for preparing a composition for treating hyperproliferative disorder, e.g., cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          antisense; PPAR-delta; hybridisation; inhibitor;
phosphorothioate linkage; 2'-O-methoxyethyl sugar; 5-methylcytosine;
hyperproliferative disorder; cancer; cytostatic; gene therapy; ds.
                                                                                                                                                                                                                                                                                                                                           Gaps
                                                                                                                                                                                                                                                                                                                                           ·,
                                                                                                                                                                                                                                                                                                                     0.8%; Score 14.2; DB 1; Length 20; 84.2%; Pred. No. 8.7e+02;
                                                                                                                                                                                                                                                                                                                                          Indels
                                                                                                                                                                                                                                                                                                 Sequence 20 BP; 5 A; 9 C; 3 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                           0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Murine PPAR-delta target site ID 137749.
                                                                                                                                       Example 16; SEQ ID NO 124; Opp; English.
                                                                                                                                                                                                                                                                                                                                                               344 TGAAGATGGGGTCTGATGG 362
                                                                                                                                                                                                                                                                                                                                                                              19 TGCAGATGGGCTGTGATGG 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                         BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                         ADL34964 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Freier SM,
                                                     Freier SM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2004-282460/26.
                                                                          WPI; 2004-282460/26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (GAAR/) GAARDE W.
(FREI/) FREIER S M.
(WAIT/) WATT A T.
                                                                                                                                                                                                                                                                                                                              Best Local Similarity
Matches 16; Conserv
        (GAAR/) GAARDE W.
(FREI/) FREIER S N
(WATT/) WATT A T.
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nucleic acid encoding PPAR delta, which specifically hybridise to and inhibit expression of PPAR-delta. The oligonucleotide specifically hybridises with at least an 8-nucleobase portion of an active site on the nucleic acid molecule encoding the PPAR-delta and comprises at least one modified internucleoside linkage, which is a phosphorothicate linkage, at least one modified sugar moiety, which is a 2'-0-methoxyethyl sugar moiety or at least one modified nucleobase, which is a 5-methylotycsine. The antisense oligonucleotides are useful for preparing a composition for
                                                                                                                                                                                                                                                                                                                                                    treating hyperproliferative disorders, e.g., cancer. The oligonucleotides of the invention have cytostatic activity and can be used for gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New oligonucleotides, which specifically hybridizes with Jun N-terminal kinase protein, useful in diagnosing or treating inflammation, fibrosis or a fibrotic or hyperproliferative disease or condition.
                                                                                                           invention describes novel antisense oligonucleotides targeted to a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Rat; Jun N-terminal kinase; JNK; Jun N-terminal kinase 1; JNK1; hyperproliferative disease; cell cycle progression; protebrorylation; tumour growth; cancer; apoptosis; prostate cancer; inflammation; fibrosis; fibrotic disease; scarring; peritoneal adhesion; lung fibrosis; conjunctival scarring; cytostatic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           .
0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Rat Jun N-terminal kinase 1 (JNK1) oligonucleotide #11.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaarde WA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 20 BP; 3 A; 3 C; 9 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 14.2; DB 1;
Pred. No. 8.7e+02;
0; Mismatches 3;
hyperproliferative disorder, e.g., cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Nero PS,
                                                     Example 16; SEQ ID NO 262; Opp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 344 TGAAGATGGGGTCTGATGG 362
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       97US-00910629.
98US-00130616.
99US-00287796.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         99US-00396902
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 84.2%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Local Similarity 84.2
nes 16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2004-168941/16.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MCKAY R.
DEAN N M.
MONIA B P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Dean NM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Rattus norvegicus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NERO P S. GAARDE W ;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          13-AUG-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      07-AUG-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             07-APR-1999;
15-SEP-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    12-FEB-2004.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                               therapy.
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(GAAR/)
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invention also relates to a pharmaceutical composition comprising the oligonucleotide (s) or its bioequivalent and a pharmaceutical carrier, a method of treating an animal having, suspected of having or prone to having a hyperproliferative disease, a method of modulating the expression of a JNK protein in cells or tissues, a method of modulating the expression of a cellular protein phosphorylation of a protein phosphorylation of a protein phosphorylation of a cellular protein that promotes one or more metastatic events in cultured cells or the cells of an animal, a method of inhibiting the growth of a tumour in an animal, a method of inducing apoptosis in a cell, a method of treating a human having a disease or condition associated with a JNK protein. The oligonucleotide and composition are useful in diagnosing or treating a disease or condition are useful in diagnosing or treating a disease or condition are useful in disease or condition (e.g. prostate cancer), a disease or condition associated with a JNK protein (e.g. inflammation, fibrosis), a fibrosis, conjunctival scarring) or a hyperproliferative disease or condition (e.g. conjunctival scarring) or a hyperproliferative disease or condition (e.g. conjunctival scarring) or a hyperproliferative disease or condition (e.g. cencer), or in inhibiting the growth of a tumour. This sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ·:
                                                                                                                                                                                                  connected by covalent linkages, where the oligonucleotide has a sequence specifically hybridisable with a nucleic acid encoding a Jun N-terminal kinase (JNK) protein and modulates the expression of the JNK protein. The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      chimeric; antisense oligonucleotide; phosphorothioate; human; microsomal prostaglandin E2 synthase; mPGES-1; mPGES-1 inhibitor; microsomal prostaglandin E2 synthase inhibitor; cytostatic; antidiabetic; immunomodulator; cardiant; neuroprotective; antiinflammatory; neuroprotective; antiinflammatory; neuroprotective; antiinflammation; immunomodulatory; cardiavascular; gene therapy; inflammation; Alzheimer's disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder; cardiovascular disorder; neurological disorder; s.
                                                                                                                                                 The invention relates to an oligonucleotide comprising 8-30 nucleotides
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
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residues are 5-methylcytidines"
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84.2%; Pred. No. 8.7e+02;
tive 0; Mismatches 3; Indels
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16. .20
/*tag= c
Example 8; SEQ ID NO 121; 71pp; English.
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nes 16; Conservative
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ADM14468/c
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The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin E2 synthase (mPGES-1). The human mEGES-1 gene is located on chromosome 9, more specifically to 9934.3. The present invention also describes: (1) antisense compounds, having a sequence comprising 8-30 bp targeted to a nucleic acid encoding mPGES-1, which specifically hybridise with the nucleic acid encoding inhibits its expression; (2) a method of inhibiting the expression of mPGES-1 in cells or tissues; and (3) a method of treating an animal having a disease or condition associated with mPGES-1. MPGES-1 chimeric antisense oligonucleotides and antisense compounds have cytostatic, antidiabetic, immunomodulator, cardiant, neuroprotective, continhimanological, immunomodulatory and cardiovascular activities, and can be used for preparing a omposition for treating a disease or condition associated with mPGES-1 e.g., inflammation, Alzheimer's condition associated with mPGES-1 e.g., inflammation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         disease, arthritis, diabetes, cancer, ischaemia or reperfusion injury, or ophthalmic, immunological, cardiovascular or neurological disorder.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  chimeric; antisense oligonucleotide; phosphorothioate; human; microsomal prostaglandin E2 synthase; mPGBS-1; mPGBS-1; inhibitor; microsomal prostaglandin E2 synthase; inhibitor; cytostatic, antidiabetic; immunomodulator; cardiant; neuroprotective; antiinflammatory; neuroprotective; antiarthritic; vasotropic; ophthalmological; immunomodulatory; cardiant; gene therapy; inflammation; Alzheimer's disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder; acadiovascular disorder; neurological disorder;
                                                                                                                                                                                                                                                                                                                                                                                                                     encoding mPGES-1, useful for preparing a composition for treating e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer or
                                                                                                                                                                                                                                                                                                                                                                                                  New antisense compound, having a sequence targeted to a nucleic acid
encoding mPGES-1, useful for preparing a composition for treating e.c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0.8%; Score 14.2; DB 1; Length 20;
34.2%; Pred. No. 8.7e+02;
ve 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 20 BP; 3 A; 8 C; 6 G; 3 T; 0 U; 0 Other;
/mod_base= OTHER
/note= "2'-0-methoxyethyls"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 4; SEQ ID NO 655; 132pp; English.
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                                                                                                                                                                 25-SEP-2003; 2003WO-US030374.
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ses 16; Conservative
                                                                                                                                                                                                                                                           (PHAA ) PHARMACIA CORP
                                                                                                                                                                                                                                                                                                                                                    WPI; 2004-305094/28.
                                                                      WO2004028458-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                        encoding mPGES-1
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                                                                                                                   08-APR-2004
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Matches
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(first entry)

ADM15006;

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microsomal prostaglanding 2 synthase; mPGES-1; mPGES-1 inhibitor; microsomal prostaglandin E2 synthase; mPGES-1; mPGES-1 inhibitor; microsomal prostaglandin E2 synthase inhibitor; cytostatic; antidiabetic; immunomodulator; cardiant; neuroprotective; antiinflammatory; neuroprotective; nootropic; antiarthritic; vasotropic; ophthalmological; immunomodulatory; cardiovascular; gene therapy; inflammation; Albahame; albabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder; ardiovascular disorder; neurological disorder; ss.
                                                                                                  Human mPGES-1 chimeric antisense oligonucleotide SEQ ID NO:1193.
                                                                                                                                              chimeric; antisense oligonucleotide; phosphorothioate; human;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             25-SEP-2003; 2003WO-US030374
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modified_base
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Synthetic.
                                                             01-JUL-2004
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The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin E2 synthase (mPGES-1). The human microsomal prostaglandin E2 synthase (mPGES-1). The human microsome is located on chromosome 9; more specifically to operate the present invention also describes: (1) antisense compounds, having a sequence comprising 8-30 bp targeted to a nucleic acid encoding meGES-1, which specifically hybridise with the nucleic acid meGES-1 and inhibits its expression; (2) a method of inhibiting the expression of mPGES-1 in cells or tissues; and (3) a method of treating an animal can anisense oligonucleotides and antisense compounds have cytostatic, antisense oligonucleotides and antisense compounds have cytostatic, antisense oligonucleotides and antisense compounds have cytostatic, antisense compounds continual antisense compounds of antishflammatory, neuroprotective, nootropic, antishflatmic, vasotropic, ophthalmological, immunomodulatory and cardiovascular activities, and can be used as mPGES-1 inhibitors and in gene therapy. The antisense compound can be used as mPGES-1 inhibitors and in gene therapy. The antisense compound contition associated with mPGES-1 e-g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer, ischemia or reperfusion injury, or condition associated with mPGES-1 e-g., inflammation reperfusion injury, or condition immunological, cardiovascular or neurological disorder.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New antisense compound, having a sequence targeted to a nucleic acid encoding mPGES-1, useful for preparing a composition for treating e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer or
                                                                                                                                                                  /note= "phosphorothioate linkages and all cytidine residues are 5-methylcytidines"
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                                                                                                                                                                                                                                                   /mod_base= OTHER
/note= "2'-0-methocyethyls"
                                                                                                                                                                                                                                                                                                                                                               /note= "2'-0-methoxyethyls"
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                                                                                   Location/Qualifiers
                                                                                                                                                   'mod base= OTHER
                                                                                                                                                                                                                                                                                                                                           mod base= OTHER
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Best Local Similarity
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                                                                                                         modified base
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                     Homo sapiens.
Synthetic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ischemia
                                                                                       Key
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/mod_base= OTHER /note= "phosphorothioate linkages and all cytidine

Location/Qualifiers

Q

'*tag=

residues are 5-methylcytidines"

/mod_base= OTHER /note= "2'-0-methocyethyls"

/*tag= a

note = "2'-0-methoxyethyls"

base= OTHER

O

16. .20 /*tag=

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antisense oligonucleotides and antisense compounds have cytostatic, antidiabetic, famunomodulator, cardiant, neuroprotective, vasotropic, antiarthritic, vasotropic, ophthalmological, immunomodulatory and cardiovascular activities, and can be used as mpGRS-1 inhibitors and in gene therapy. The antisense compound can be used for preparing a composition for treating a disease or condition associated with mpGES-1 e.g., inflammation, Alzheimer's
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin E2 synthase (mPGES-1). The human mPGES-1 gene is located on chromosome 9, more specifically to 9334.3. The present invention also describes: (1) antisense compounds, having a sequence comprising 8-30 bp targeted to a nucleic acid encoding mPGES-1, which specifically hybridise with the nucleic acid encoding inhibits its expression; (2) a method of inhibiting the expression of mPGES-1 in cells or tissues; and (3) a method of treating an animal having a disease or condition associated with mPGES-1. MPGES-1 chimeric
                                                                                                      New antisense compound, having a sequence targeted to a nucleic acid encoding mPGES-1, useful for preparing a composition for treating e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer or
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                                                                                                                                                                                                                                                                                                                                                                                                      Claim 4; SEQ ID NO 1193; 132pp; English.
WPI; 2004-305094/28.
                                                                                                                                                                                                                                                                                                      ischemia.
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Gaps

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Indels

0; Mismatches

1433 CAGAGGATGCCATGAAACA 1451

Conservative

16;

Matches

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CCGAGGATGCCCTGAGACA

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g 8

ADM15006 standard; DNA; 20 BP.

RESULT 1144

ADM15006

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06-MAY-2004.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ņ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The invention relates to antisense oligonucleotides targeted to cyclin-
                                                                                                                             Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /note= "Phosphorothioate backbone. All cytidines are
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New compounds, particularly antisense oligonucleotides targeted to a nucleic acid encoding cyclin-dependent kinase 6, useful for treating cancer, bacterial/viral infection or conditions involving aberrant
ophthalmic, immunological, cardiovascular or neurological disorder.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               nucleotides"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /mod_base= OTHER
/note= "2'-methoxyethyl (2'-MOE) nucleotides"
                                                                               0.8%; Score 14.2; DB 1; Length 20;
84.2%; Pred. No. 8.7e+02;
ative 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                               Cyclin-dependent kinase 6, antisense oligonucleotide #153
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            infection;
                                       Sequence 20 BP; 2 A; 6 C; 7 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /mod_base= OTHER
/note= "2'-methoxyethyl (2'-MOE)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    antisense therapy; cyclin-dependent kinase 6; hyperproliferative disorder; cancer; bacterial viral infection; apoptosis; ss; probe; human.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Location/Qualifiers
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                                                                                                                                                                  1433 CAGAGGATGCCATGAAACA 1451
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                                                                                                                                                                                                                                                                                                                    ADO56089 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                     29-JUL-2004 (first entry)
                                                                    Query Match
Best Local Similarity 84.2
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'*tag=
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Dobie KW;
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modified_base
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                                                                                                                                                                                                                                                                                                                                                             ADO56089;
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dependent kinase 6 antisense oligonucleotide. Note: Segid 15-134 are also used in the sequence listing but these sequences do not match segid 15-134 displayed in Tables 1 and 2 (page 30-34).
                                                                                                                                                                                                                                                                                            Gaps
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                                                                                                                                                                                                             Query Match 0.8%; Score 14.2; DB 1; Length 20; Best Local Similarity 84.2%; Pred. No. 8.7e+02; Matches 16; Conservative 0; Mismatches 3; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Cyclin-dependent kinase 6, antisense oligonucleotide #220
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        antisense therapy, cyclin-dependent kinase 6; hyperproliferative disorder, cancer, bacterial infection, viral infection, apoptosis, ss; probe, human.
                                                                                                                                          Sequence 20 BP; 7 A; 5 C; 5 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 20 BP; 3 A; 5 C; 5 G; 7 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                            923 TGTTCCAGCTGCTCGTGG 941
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                                                                                                                                                                                                                                                                                                                                                                                                     19 TGTTTCAGCTTCTCCGAGG 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               31-JUL-2002; 2002US-00210802.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Dobie KW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2004-356241/33.
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The invention relates to antisense oligonucleotides targeted to cyclindependent kinase 6, and which inhibit the expression of cyclindependent kinase 6. The antisense oligonucleotides are useful for treating a disease or condition associated with cyclin-dependent kinase 6, such as a hyperproliferative disorder (e.g. cancer), or conditions arising from pacterial or viral infections, or involving aberrant apoptosis. They are also useful in research and diagnostics for modulating the expression of cyclin-dependent kinase 6. The present sequence represents a cyclindependent kinase 6 antisense oligonucleotide. Note: Seqid 15-134 are also used in the sequence listing but these sequences do not match seqid 15-134 displayed in Tables 1 and 2 (page 30-34).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /note= "Phosphorothioate backbone. All cytidines are
methylcytidines."
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mod_base= OTHER
/note= "2'-methoxyethyl (2'-MOE) nucleotides"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     'note= "2'-methoxyethyl (2'-MOE) nucleotides"
                                                                                                                                                                                                                                               Cyclin-dependent kinase 6, antisense oligonucleotide #154
                                                                                                                                                                                                                                                                                                  antisense therapy; cyclin-dependent kinase 6;
hyperproliferative disorder; cancer; bacterial infection;
viral infection; apoptosis; ss; probe; human.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Location/Qualifiers
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/mod_base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /mod_base= OTHER
                                                             ADO56090 standard; DNA; 20 BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           modified base
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           apoptosis.
                                                                                                                       ADO56090;
RESULT 1147
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                                ADO56090/
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The invention relates to a compound targeted to a nucleic acid molecule encoding the human PIM-1 polypeptide. The compound is an antisense oligomucleotide that specifically hybridises with the mucleic acid and inhibits expression of the polypeptide. The antisense oligomucleotide comprises at least one modified internucleoside linkage i.e. a phosphorothioate linkage, at least one modified sugar moiety, preferably a 2'-0-methoxyethyl sugar moiety, or at least one modified nucleobase comprising a 5-methyloryosine. The antisense compounds are useful for modularing the expression of the human PIM-1 polypeptide and in preparation of a composition for treating hyperproliferative disorders, e.g. cancer. This sequence repersents a human PIM-1 DNA antisense oligonucleotide of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New compound that modulates PIM-1 expression, useful in treating an animal having a disease or condition, i.e. hyperproliferative disorder.
                                                                                                                                                                                                                   Human; PIM-1; ss; antisense oligonucleotide; phosphorothioate linkage;
2'-0-methoxyethyl sugar moiety; 5-methylcytosine;
hyperproliferative disorder; cancer; cytostatic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0.8%; Score 14.2; DB 1; Length 20; 44.2%; Pred. No. 8.7e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 20 BP; 3 A; 6 C; 8 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                         Human PIM-1 DNA antisense oligonucleotide #24.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 15; SEQ ID NO 36; 51pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        966 GGTGCTACACCGAGACCTC 984
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CAGCTTCTCCGAGGTCTGG 1
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                                                                                                                                                           (first entry)
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                                                                                               ADN03067 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2004-374981/35.
                                                                                                                                                                                                                                                                                                                                                                                                                                               (ISIS-) ISIS PHARM
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                                                                                                                                                                                                                                                                                                                     US2004092463-A1
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                   19
                                                                                                                              ADN03067;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Watt AT;
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                                                                 RESULT 1148
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                                                                                 ADN03067,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADN0313
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Gaps

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Indels

0.8%; Score 14.2; DB 1; Length 20; 84.2%; Pred. No. 8.7e+02;

0; Mismatches

Matches 16; Conservative

Query Match Best Local Similarity

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The invention relates to a compound targeted to a nucleic acid molecule encoding the human PIM-1 polypeptide. The compound is an antisense oligonucleotide that specifically hybridases with the nucleic acid and inhibits expression of the polypeptide. The antisense oligonucleotide comprises at least one modified internucleoside linkage i.e. a posphorothhoate linkage, at least one modified sugar moiety, preferably a 2'-O-methoxyethyl sugar moiety, or at least one modified nucleobase comprising a 5-methylcytosine. The antisense compounds are useful for modulating the expression of the human PIM-1 polypeptide and in preparation of a composition for treating hyperproliferative disorders, e.g. cancer. This sequence repersents a human PIM-1 DNA antisense oligonucleotide target region of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       detection; fungal infection; soil fungal infection; vegetable fungal infection; pathogenic fungus; Microcentrospora acerina; Fibularhizoctonia carotae; Pythium; hybridisation; primer; ss.
                                                                                                                                                                                                                                                                                                                                          New compound that modulates PIM-1 expression, useful in treating an animal having a disease or condition, i.e. hyperproliferative disorder.
Human; PIM-1; ss; antisense oligonucleotide; phosphorothioate linkage;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0.8%; Score 14.2; DB 1; Length 20; 84.2%; Pred. No. 8.7e+02; ative 0; Mismatches 3; Indels
                 2'-0-methoxyethyl sugar moiety; 5-methylcytosine;
hyperproliferative disorder; cancer; cytostatic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 20 BP; 3 A; 8 C; 6 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                               Example 15; SEQ ID NO 103; 51pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   966 GGTGCTACACCGAGACCTC 984
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   20
                                                                                                                                                                      11-NOV-2002; 2002US-00292849.
                                                                                                                                                                                                        11-NOV-2002; 2002US-00292849.
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01-NOV-2002; 2002GB-00025551.
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Best Local Similarity 84.2
Matches 16, Conservative
                                                                                                                                                                                                                                         (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                                                                           WPI; 2004-374981/35.
                                                                                                  US2004092463-A1
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                                                                     Homo sapiens
                                                                                                                                    13-MAY-2004
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The present invention describes an assay for detecting fungal infection of soil or vegetables by pathogenic fungal species, particularly Microcentrospora acerina, Fibularhizoctomia carotae or Pythium species. The assay comprises: (1) obtaining a sample of soil or vegetable; (2) treating the sample to lyse fungal cells; (3) effecting a PCR on DNA celeating DNA fragments generated by the PCR. Also pair; and (4) detecting DNA fragments generated by the PCR. Also described: (1) an 18-24-mer oligonucleotide primer by described: (1) an 18-24-mer oligonucleotide primer; (3) a substrate oligonucleotide sequence selected from SEQ ID NO:1 to 28; (2) a substrate oligonucleotide and mobblised on it the 18-24-mer oligonucleotide primer; (3) a primer composition comprising a pair of 18-24-mer oligonucleotide primers; (4) a kit for performing the assay method, for nucleic acid extraction from host vegetable tissue. The assay is useful for detecting fungal correction of soil or vegetables by pathogenic fungal species, infection of soil or vegetables by pathogenic fungal species. The present sequence represents a general primer which is given in the exemplification of the present invention.
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                                                                                                                                                               Detecting fungal infection of soil or vegetables by Microcentrospora acerina, Fibularhizoctonia carotae or Pythium species by treating the sample of soil or vegetable and effecting a PCR on DNA released by lysis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ·:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0.8%; Score 14.2; DB 1; Length 20;
34.2%; Pred. No. 8.7e+02;
ve 0; Mismatches 3; Indels
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                                                                                      Lund
                                                                                   Wanner L,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 20 BP; 2 A; 6 C; 5 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                           Disclosure; SEQ ID NO 30; 44pp; English.
                                                                                   Naerstad R,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1549 CTTCGGTCTTCGTCGATGC 1567
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    84.2%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Best Local Similarity 84.2
Matches 16; Conservative
                 (COCK/) COCKBAIN J R M.
                                                                                   Klemsdal
                                                                                                                                                                                                                               of the fungal cells.
                                                                                                                           WPI; 2004-376207/35.
                                                                                   Hermansen A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
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20 CTGCGTTCTTCATCGATGC N qq

.; 0

Gaps 0;

detection; fungal infection; soil fungal infection; vegetable fungal infection; pathogenic fungus; Microcentrospora acerina; Fibularhizoctonia carotae; Pythium; hybridisation; primer; ss. Pungi, oomycete and plant general hybridisation primer C SEQ ID NO:31. ВP ADN61577 standard; DNA; 20 (first entry) 29-JUL-2004 Synthetic. ADN61577; RESULT 1151 ADN61577,

31-OCT-2003; 2003WO-GB004712. 01-NOV-2002; 2002GB-00025550.

WO2004040017-A2.

13-MAY-2004.

(CARR-) CARROTECH AS

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The present invention describes an assay for detecting fungal infection of soil or vegetables by pathogenic fungal species, particularly conscient of species, particularly secretar or pertain species.

Microcentrospora acerina, Fibularhizoctonia carctae or Pythium species. The assay comprises: (1) obtaining a sample of soil or vegetable; (2) treating the sample to lyse fungal cells; (3) effecting a PCR on DNA treating DNA fragments generated by the PCR. Also described: (1) an 18-24-mer oligomucleotide primer and (4) detecting DNA fragments generated by the PCR. Also oligomucleotide sequence selected from SEQ ID NO:1 to 28; (2) a substrate conjugant inmobilised on it the 18-24-mer oligomucleotide primer; (3) at primer composition comprising a pair of 18-24-mer oligomucleotide primer; (3) a kit for performing the assay method, for nucleic acid extraction from host vegetable tissue. The assay is useful for detecting fungal infection of soil or vegetables by pathogenic fungal species. The present sequence represents a general primer which can hybridise to DNA from all fungi, all compretes and all plants, which
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ö
                                                                                            Detecting fungal infection of soil or vegetables by Microcentrospora acerina, Fibularhizoctonia carotae or Pythium species by treating the sample of soil or vegetable and effecting a PCR on DNA released by lysis of the fungal cells.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Glutamine-fructose-6-phosphate amidotransferase; diabetes; ischemia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         .
0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0.8%; Score 14.2; DB 1; Length 20; 84.2%; Pred. No. 8.7e+02; ive 0; Mismatches 3; Indels
                              Lund G
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    given in the exemplification of the present invention.
                              Wanner L,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 20 BP; 7 A; 5 C; 6 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Chimeric phosphorothioate oligonucleotide #343
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /mod_base= other
/note= "2-methoxyethyl wing"
17. .20
/*tag= b
/mod_base= other
/note= "2-methoxyethyl wing"
                                                                                                                                                                                      Disclosure; SEQ ID NO 31; 44pp; English.
                              Naerstad R,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match 0.8
Best Local Similarity 84.2
Matches 16; Conservative
                            Hermansen A, Klemsdal S,
(COCK/) COCKBAIN J R M.
                                                                 WPI; 2004-376207/35.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                reperfusion; ss.
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modified_base
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADP76544;
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                                                                                                                                                                            New compounds, particularly antisense oligonucleotides targeted to a nucleic acid encoding glutamine-fructose-6-phosphate amidotransferase (GFAI), for treating diabetes, a cardiovascular or neurologic disorder, ischemia/reperfusion injury.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Glutamine-fructose-6-phosphate amidotransferase; diabetes; ischemia;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 20 BP; 4 A; 6 C; 2 G; 8 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                          oligonucleotides inhibit human GFAT expression.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Chimeric phosphorothioate oligonucleotide #149.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /mod_base= other
/note= "2-methoxyethyl wing"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /note= "2-methoxyethyl wing"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0; Mismatches
                                                                                                                                                                                                                                                         Claim 4; SEQ ID NO 343; 175pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Location/Qualifiers
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/mod_base= other
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  131 GGATGAAGAAGATCAAACG 149
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                19 GGATGAAGAAGTTCACAAG 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADP76350 standard; DNA; 20 BP
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                            02-OCT-2003; 2003WO-US03332.
                                                         17-OCT-2002; 2002US-0419268P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      84.2%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Local Similarity 84.2
les 16; Conservative
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                                                                                                                      Crosby SD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /*tag=
                                                                                      (PHAA ) PHARMACIA CORP
                                                                                                                                                 WPI; 2004-348453/32.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    reperfusion; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Key
modified_base
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                                                                                                                      Broschat KO,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               29-APR-2004
29-APR-2004.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADP76350;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
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The present invention relates to a compound which specifically hybridizes with a nucleic acid molecule encoding GFAT, and inhibits the expression of GFAT. Specifically claimed are antisense oligonucleotides capable of modulating the expression of GFAT, and which comprise any of the 30G3 sequences of 20 base pairs, given in the specification. The compound, composition and methods are useful for treating a disease or condition associated with GFAT, such as a disease or condition, e.g. diabetes, a cardiovascular or neurological disorder, ischemia/reperfusion injury. They are also useful in research and diagnostics for modulating the expression of GFAT. The present sequence represents a chimeric phosphorothicate oligonucleotide with 2'-MOE wings and a decoyy gap, these oligonucleotides inhibit human GFAT expression.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SF-25 antigen; magnetic bead; cancer; cancer diagnosis; PCR; primer; 4F2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Examining cancer cells through isolation of such cells expressing SF-25 antigen on cell surface for bonding to magnetic beads with antigen-
                                                                                                                                                                    New compounds, particularly antisense oligonucleotides targeted to a nucleic acid encoding glutamine-fructose-6-phosphate amidotransferase (GFAT), for treating diabetes, a cardiovascular or neurologic disorder, ischemia/reperfusion injury.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0.8%; Score 14.2; DB 1; Length 20;
84.2%; Pred. No. 8.7e+02;
tive 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 20 BP; 3 A; 6 C; 3 G; B T; 0 U; 0 Other;
                                                                                                                                                                                                                                                    Claim 4; SEQ ID NO 149; 175pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mitsunaga
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 4F2 gene measuring reverse primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           131 GGATGAAGAAGATCAAACG 149
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0
                                 17-OCT-2002; 2002US-0419268P,
 02-OCT-2003; 2003WO-US03332.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GGATGAAGAAGTTCACAAG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADO44487 standard; DNA; 20
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Best Local Similarity 84.2
Matches 16; Conservative
                                                                                                   Broschat KO, Crosby SD;
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                                                                (PHAA ) PHARMACIA CORP.
                                                                                                                                    WPI; 2004-348453/32.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2004-419769/39
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WO2004042401-A1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 12-AUG-2004
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                                                                                                                                                                                                                                                                                                                                            4F2
                                                                                                     The invention relates to detecting cancer cells and involves isolating cancer cells expressing SP-25 antigen on cell surface from a living body for bonding to magnetic beads with use of antigen-antibody reaction of the cancer cell and anti SP-25 antibody or its antigen-binding fragment, collecting the beads and examining the bound cancer cells. The method is collecting the beads and examining the bound cancer cells. The method is including leukemia. The method is convenient and efficient, without needing any special equipment like cell sorrers. The present sequence represents a PCR primer used in measuring the expression level of the 4F?
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /mod_base= OTHER
/note= "OTHER = Phosphorothioate backbone, bases 1-5 and
16.20 2'. MOE wing bases, all cytidine residues are 5-
methylcytidines"
  antibody reaction, applicable in diagnosis of cancer including leukemia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Antisense 2'-MOE gapmer oligo targeted to human ApoB RNA - SEQ 420
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0
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84.2%; Pred. No. 8.7e+02;
iive 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    phosphorothioate backbone; human; chromosome 2p23-2p24; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 20 BP; 5 A; 8 C; 3 G; 4 T; 0 U; 0 Other;
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                                                 Example 4; SEQ ID NO 16; 33pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              208 GAGCAGATAGGCCTGGATG 226
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2003WO-US015493
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Local Similarity
es 16; Conserv
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15-MAY-2003;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       12-AUG-2004
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AD032972;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Matches
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Gaps

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The invention relates to a novel antisense compound where the compound hybridises to and inhibits expression of mRNA encoding human apolipoprotein B (ApoB) after 16-24 hours by at least 18% in 80% confound the invention demonstrates cardiovascular, antiated compound of the invention demonstrates cardiovascular, antiated compound of the invention demonstrates cardiovascular, antiated compound of the invention demonstrates cardiovascular, compound of the invention demonstrates cardiovascular, cardiant, vasotropic, national catholic, eating disorder-related, cytostatic, candocrine, vasotropic, neuroprotective and nootropic activities and may consocrated for inhibiting the expression of apolipoprotein B in cells or tissues in vivo in order to address a condition associated with abnormal lipid or cholesterol metabolism. The compound may be useful for cholesterol metabolism. The compound may be useful for cholesterol levels, lipid levels, fatty acid levels, acute phase cholesterol and chylomicrons and thus may be utilised during treatment of reactants and chylomicrons and thus may be utilised during treatment of syndrome, sexual ateliotic dwarfism, hyperthyloxidism, hypertension, cardiovascular disorders, Von Gierke's disease, lipodystrophy, componence, obstructive liver disease, Altheimer's disease, dementia, diabetes, obesity and atherosclerosis. The current sequence is that of an expense 2'-now Dayn Dayn Dayn Dayner 
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 apolipoprotein B; ApoB; cardiovascular; antiarteriosclerotic; antilipaemic; antidiabetic; anorectic; cardiant; vasotropic; hypotensive; anabolic; eating disorder; cytostatic; endocrine; vasotropic; neuroprotective; nootropic; lipid; cholesterol metabolism; hyperlipoproteinaemia; hyperlipidsemia; hypercholesterolaemia; von Gierke's disease; lipodystrophy; Cushing's syndrome; sexual ateliotic dwarfism; hyperthyroidism; hypertension; anorexia nervosa; Werner's syndrome; hepatcoma; multiple myeloma; uraemia; impotence; obstructive liver disease; Alzheimer's; dementia; diabetes; obesity; atherosclerosis; antisense; 2'-MOE gapmer; 2'-methoxyethyl wing;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /note= "OTHER = Phosphorothioate backbone, bases 1-5 and 16-20 2'-MOB wing bases, all cytidine residues are 5-methylcytidines"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
Antisense oligonuclectide compound that inhibits expression of mRNA encoding human apolipoprotein B, useful for treating hyperlipidemia, diabetes, obesity, von Gierke's disease, lipodystrophies, Cushing's
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Antisense 2'-MOE gapmer oligo targeted to human ApoB RNA - SEQ 128
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 phosphorothioate backbone; human; chromosome 2p23-2p24; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 20 BP; 9 A; 5 C; 4 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0; Mismatches
                                                                                                                                             Example 33; SEQ ID NO 420; 483pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                583 CTATCTGAGATTGGCTTTG 601
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               19 CTTTCTCAGATTGGCTTGG 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADO32680 standard; DNA; 20 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ilarity 84.2%;
Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            targeted to human ApoB RNA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
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les 16; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Key
modified_base
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
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Matches
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The invention relates to a novel antisense compound where the compound hybridises to and inhibits expression of mRNA encoding human apolipoprotein B (ApoB) after 16-24 hours by at least 30% in 80% confluent HepG2 cells in culture at a concentration of 150 nM. The compound of the invention demonstrates cardiovascular, ancretics. Cardiant, anitateriosclerotic, antilipaemic, antidiabettc, anorectic, cardiant, carotropic, hypotensive, anabolic, eating disorder-related, cytostatic, cordine, vasotropic, neuroprotective and nootropic activities and may be useful for inhibiting the expression of apolipoprotein B in cells or citissues in vivo in order to address a condition associated with abnormal cordinates in vivo in order to address a condition associated with abnormal cordinates in vivo in order to address a condition associated with abnormal cordinates in vivo in order to address a condition associated with abnormal cordinates of inpid or cholesterol metabolism. The compound may be useful for decreasing circulating lipoprotein levels, triglycelide during treatment of hyperlipoproteinaemia, hyperlipodystrophy, Cushing's cordinovacular disorders, von Gierke's disease, lipodystrophy, Cushing's syndrome, sexual ateliotic dwarfism, hyperthyroidism, hypertenaion, cordiabetes, obesity and atherosclerosis. The current sequence is that of antisense 2'-MOE (2'-methoxyethyl) gapmer oligo of the invention which is trageted to human ApoB RNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         apolipoprotein B; ApoB; cardiovascular; antiarteriosclerotic; antilipaemic; antidiabetic; anorectic; cardiant; vasotropic; hypotensive; anabolic; eating disorder; cytostatic; endocrine; vasotropic; hypotentorotective; nootropic; lipid; cholesterol metabolism; hyporlipoproteinaemia; hyperlipidaemia; hyperlipidaemia; hypersthopy; Cushing's syndrome; sexual ateliotic dwarfism; hyperthyroidism; hypertension;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human apolipoprotein B (ApoB) antisense therapy target DNA - SEQ 517.
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                                                                                                                                                                                                                                                                                                               Antisense oligomucleotide compound that inhibits expression of mRNA encoding human apolipoprotein B, useful for treating hyperlipidemia, diabetes, obesity, von Gierke's disease, lipodystrophies, Cushing's
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84.2%; Pred. No. 8.7e+02;
tive 0; Mismatches 3; Indels
                                                                                                                                                                                                                                  Dobie KW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 20 BP; 4 A; 7 C; 4 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                  Lemonidis-Tarbet K,
                                                                                                                                                                                                                                                                                                                                                                                                                         Example 29; SEQ ID NO 128; 483pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1012 AGGGAGAGCTCAAGCTGG 1030
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AGGTATGAGCTCAAGCTGG 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADO33069 standard; DNA; 20 BP.
                                                                                                                         13-NOV-2002; 2002US-0426234P.
15-MAY-2003; 2003WO-US015493.
                                                                               13-NOV-2003; 2003WO-US036411
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                                                                                                                                                                                            (ISIS-) ISIS PHARM INC.
                                                                                                                                                                                                                                      Graham M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Local Similarity
                                                                                                                                                                                                                                                                              WPI; 2004-420321/39.
WO2004044181-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  12-AUG-2004
                                          27-MAY-2004.
                                                                                                                                                                                                                                      Crooke R,
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The invention relates to a novel antisense compound where the compound hybridises to and inhibits expression of mRNA encoding human appolipation in (Appoliter) (1-24 hours by at least 30% in 80% confluent HepGZ cells in culture at a concentration of 150 nM. The comfound of the invention demonstrates cardiovascular, confluent HepGZ cells in culture at a concentration of 150 nM. The compound of the invention demonstrates cardiovascular, confluent HepGZ cells in culture at a concentration of 150 nM. The compound of the invention demonstrates cardiovascular, cardivities and may be useful for inhibiting the expression of apolipoprotein B in cells or tissues in vivo in order to address a condition associated with abnormal lipt or cholesterol metabolism. The compound may be useful for cholesterol metabolism. Input levels, triglyceride levels, cholesterol levels, lipid levels, fatty acid levels, acute phase creations and chus may be utilised during treatment of hyperlipoproteinaemia, hyperlipidaemia, hypercholesterolaemia, cardiovascular disorders, won Giarke's disease, lipodystrophy, Cushing's syndrome, sexual ateliotic dwarfism, hyperthyroidism, hypertension, cardiovascular disease, dasease, damentia, chapters, obesity and atherosclerosis. The current sequence is that of a human apolipoprotein B (ApoB) antisense therapy target DNA of the human ApoB gene is located at chromosome 2p23-2p24.
anorexia nervosa; Werner's syndrome; hepatoma; multiple myeloma; uraemia; impotence; obstructive liver disease; Alzheimer's; dementia; diabetes; obesity; atherosclerosis; human; chromosome 2p23-2p24; ds;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human DR1-associated protein 1 antisense oligonucleotide ISIS #171285.
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Antisense oligonucleotide compound that inhibits expression of mRNA encoding human apolipoprotein B, useful for treating hyperlipidemia, diabetes, obesity, von Gierke's disease, lipodystrophies, Cushing's
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
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84.2%; Pred. No. 8.7e+02;
tive 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                    Dobie KW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 20 BP; 5 A; 4 C; 7 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                 Lemonidis-Tarbet K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example 36; SEQ ID NO 517; 483pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1012 AGGGGAGAGCTCAAGCTGG 1030
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                                                                                                                                                                                                                              13-NOV-2003; 2003WO-US036411.
                                                                                                                                                                                                                                                                          13-NOV-2002; 2002US-0426234P. 15-MAY-2003; 2003WO-US015493.
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                                                                                                                                                                                                                                                                                                                                        (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                 Graham M,
                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2004-420321/39
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Best Local Similarity
Matches 16; Conserv
                                                               antisense target
                                                                                                                                            WO2004044181-A2
                                                                                                       Homo sapiens
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                                                                                                                                                                                      27-MAY-2004
                                                                                                                                                                                                                                                                                                                                                                               Crooke R,
                                     obesity;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   syndrome
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/note= "Phosphorothioate backbone where all cytidines are 5-methyl cytidines"  
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New compounds, particularly oligonuclectides targeted to a nucleic acid encoding DR1-associated protein 1, useful for treating diseases associated with DR1-associated protein 1, e.g. developmental disorders.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The present sequence is directed to antisense oligonucleotides targeted to DRL-associated protein 1 (also known as DRAPL and negative cofactor (alpha (NC2-alpha)) and which modulates to the expression of DRL-associated protein 1. The invention is useful for treating a disease or condition associated with DRL-associated protein 1 such as developmental disorder. The present sequence is human DRL-associated protein 1 antisense oligonucleotide. This sequence is used in the
DR1-associated protein 1; DRAP1; negative cofactor 2 alpha; NC2-alpha; developmental disorder; therapy; human; antisense; phosphorothioate backbone; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    nucleic acid amplification; hybridisation; microarray; diagnosis;
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0
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34.2%; Pred. No. 8.7e+02;
ive 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Nucleic acid amplification method related competitor DNA #1.
                                                                                                                                                                                                                     -methoxyethyl nucleotides"
                                                                                                                                                                                                                                                                          -methoxyethyl nucleotides"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 20 BP; 6 A; 6 C; 4 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example 15; SEQ ID NO 16; 33pp; English
                                                                                              ocation/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 exemplification of the invention.
                                                                                                                                                                                                                                               /*tag= c
/mod_base= OTHER
/note= "2' -metho
                                                                                                                                     'mod_base= OTHER
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16. .20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           64
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                                                                                                                                                                                                                                                                                                                                                         10-DEC-2002; 2002US-00317279.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GGACCAGCAGTGTGACTGC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GGAGCAGCTTTGACTTC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                84.2%;
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                                                                                                                      Д
                                                                                                                                                                                             *tag≖ a
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                                                                                                          ...20
*tag=
                                                                                                                                                                                                                                                                                                                                                                                                                 (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                            Dobie KW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2004-440383/41.
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                                                                                             Key
modified_base
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                                                      Homo sapiens
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                                                                   Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                            Chiang M,
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                                                                                                                                                                                                                                                                                                                                                        The invention describes the efficient amplification of at least one template nucleic acid (A) comprising PCR amplification, in the presence, from the start, of a competitor (I) that inhibits formation of one of the template strands amplified by PCR. Also described is a method for cemplate strands amplified by PCR. Also described is a method for cemplate strands amplified by PCR. Also described is a method for convex then detecting the amplification product by hybridisation or with a complementary probe. The method is used to amplify nucleic acids for subsequent detection by hybridisation, particularly in a microarray format, e.g. for diagnosis, especially genetic analysis; analysis of single-nucleotide polymorphisms; detection of microarganisms and/or of viral pathogens It can also be used for real-time quantitative PCR (by cyclic repetition of the amplification and hybridisation steps). The method provides quantitative, microarray based analysis; involves fewer crept steps (amplification and detection are done as a continuous process) and has a lower error rate. The process can be done in a closed vessel and controdies a small, easily handled, system for point-of-care diagnosis. Although (I) reduces the amount of amplification product formed, it increases the hybridisation signal. This sequence represents an objection accordance of the
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                                                                                                                                                                                                                                                                          Efficient amplification of template nucleic acid, useful e.g. for genetic analysis or detecting microorganisms, uses a competitor that inhibits amplification of one template strand.
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84.2%; Pred. No. 8.7e+02;
tive 0; Mismatches 3; Indels
genetic analysis; single-nucleotide polymorphism analysis; microorganism detection; viral pathogen detection; real-time quantitative PCR; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Nucleic acid amplification method associated primer #5.
                                                                                                                                                                                                                          Bickel R, Ellinger T, Wagenhaus A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 20 BP; 7 A; 5 C; 6 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                              (CLON-) CLONDIAG CHIP TECHNOLOGIES GMBH.
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                                                                                                                                                                     19-NOV-2002; 2002DE-01053966.
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Best Local Similarity 84.2
Matches 16; Conservative
                                                                                                                                                                                                                                                   WPI; 2004-420638/39.
                                                                                    WO2004046378-A2.
                                                                                                                                                                                                                           Ermantraut E,
                                                          Unidentified.
                                                                                                               03-JUN-2004.
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The invention describes the efficient amplification of at least one template nucleic acid (A) comprising PCR amplification, in the presence, from the start, of a comperitor (I) that inhibits formation of one of the template strands amplified by PCR. Also described is a method for template strands amplified by PCR. Also described is a method for new process then detecting the amplification product by hybridisation with a complementary probe. The method is used to amplify nucleic acids for subsequent detection by hybridisation, particularly in a microarray formal pathogens. It can also be used for real-time quantitative PCR (by viral pathogens. It can also be used for real-time quantitative PCR (by cyclic repetition of the amplification and hybridisation steps). The method provides quantitative, microarray-based analysis involves fewer steps (amplification and detection are done as a continuous process) and has a lower error rate. The process can be done in a closed vessel and provides a small, easily handled, system for point-of-care diagnosis.

Although (I) reduces the amount of amplification product formed, it increases the hybridisation signal. This sequence repersents a primer of increases the hybridisation signal.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 increases the hybridisation signal. This sequence represents a primer associated with the amplification method of the invention.
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                                                                                                                                                                                                                                                                                                                                                                                                               Wagenhaus A;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 2004-420638/39
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WO2004046378-A2
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92US-00987132.
92US-00989848.
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                          89US-00399874.
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                                                                                        (GEHO ) GEN HOSPITAL CORP.
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                                                                                                            HOECHST AG
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                          29-AUG-1989;
28-AUG-1990;
                                                                                                                                                        Stengelin S,
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07-DEC-1992;
07-DEC-1992;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              pharmaceutical compositions exhibit cytostatic and antiasthmatic activities. Furthermore, it provides a transgenic animal comprising the asthma locus-1 (AST-1) DNA. This oligonucleotide sequence is a PCR primer used to sequence the exon and exon/ intron boundaries of human GPRA DNA, given in table 5 of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       locus AST-1 for asthma and other ign mediated diseases mapped to the human chromosome 7p14-p15. Specifically, it refers to two overlapping genes namely GPRA (G-protein coupled receptor for asthma susceptibility) and AAA1 (asthma associated alternatively spliced gene 1). The present invention describes identifying single nucleotide polymorphisms, as well as insertion or deletion polymorphisms, occurring at different positions in the AST-1 locus, and furthermore providing vectors, host cells, primers and probes in order to determine the status of an individual. Accordingly, it provides a kit to diagnose or assess predisposition to asthma, chronic obstructive pulmonary disease or cancer and other IgE mediated diseases including rhinitis and dermatitis, such that derived
                                                                                                                                                                                                                                                                                                                                       New GPRA polypeptides, useful in preparing a composition for diagnosing, treating or preventing asthma, other IgE-mediated disease, chronic obstructive pulmonary disease or cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      This invention relates to the identification of a novel susceptibility
                                                                                                                                                                                                                                        Vendelin J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Encodes ballast constituent in pINT69d pro-insulin fusion protein.
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                                                                                                                                                                                                                                        Polvi A, Maekelae S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 20 BP; 4 A; 6 C; 4 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                        Laitinen LA,
                                                                                                                                                                                                                                                                                                                                                                                                                              Example 7; Page 76; 265pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        210 GCAGATAGGCCTGGATGAG 228
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GCAAATATCCCTGGATGAG 2
                                                                                 20-DEC-2002; 2002US-0435846P.
03-JAN-2003; 2003US-0437895P.
26-MAR-2003; 2003US-0458767P.
09-JUL-2003; 2003US-0466000P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          BP
                                          19-DEC-2003; 2003WO-FI000973.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    92US-00838221.
                                                                                                                                                                                                                                                          Salmikangas P;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAQ51806 standard; DNA; 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Local Similarity 84.2
les 16; Conservative
                                                                                                                                                                                                                                      Kere J,
                                                                                                                                                                                                                                                                                                  WPI; 2004-500286/47.
                                                                                                                                                                                              (GENE-) GENEOS OY.
                                                                                                                                                                                                                               Laitinen T,
Pulkkinen V,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    23-APR-1992;
  08-JUL-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            20-DEC-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                US5227293-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        13-JUL-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAQ51806;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RESULT 1162
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Best Loca
Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAQ51806
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Prepn. of fusion proteins contg. ballast constituent and protein - giving
                                                                                                                                                                                                                                                                                                                                                                                                  Sequence AAQ51806 is a specific example of the novel generic ballast constituent coding sequence. The invention covers fusion proteins in which a short ballast constituent is fused to a desired protein, esp. to modified pro-insulin, to increase recombinant production of the protein. See AAQ51798-Q51799 and AAQ51802-Q51811
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   inhibitor; inhibition; ribozyme; treatment; prevention; psoriasis; asthma; inflammatory diseases; restenosis; cardiovascular condition;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Enzymatic RNA molecules which cleave mRNA - used to treat or prevent inflammatory, arthritic, stenotic or cardiovascular diseases or conditions.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Specific; cleavage; target RNA; protein; prophylaxis; expression;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       .;
0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match 0.8%; Score 14.2; DB 1; Length 21; Best Local Similarity 84.2%; Pred. No. 9.1e+02; Matches 16; Conservative 0; Mismatches 3; Indels
Seed B;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 21 BP; 10 A; 6 C; 3 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Enzymatic RNA molecule c-myb mRNA target sequence.
                                                                                                                                                                                                                                        prods. which are protease resistant or insoluble
Uhlmann E,
Habermann P,
                                                                                                                                                                                                                                                                                                                     Example 8; Col 7-8; 22pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            885 TGGGAACATCATCAACATG 903
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2 TGGCAACACATCAACACG 20
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A new method for the detection of nucleic acids comprises (a) amplifying a DNA by PCR using primers to which an appropriate RNA polymerase promoter has been appended; (b) transcribing the amplified DNA into RNA using an RNA polymerase; (c) forming RNA:DNA hybrids; and (d) immunochemically detecting the RNA:DNA hybrids using antibodies directed to RNA:DNA hybrids using antibodies directed to RNA:DNA hybrids using antibodies directed the RNA:DNA hybrids using antibodies directed the RNA:DNA hybrids using antibodies directed from the hlyA gene and spanned a 730 base pair region of the gene from nucleotides 602-1332. Por further use in the invention, the primer described in AAT42247 had an additional 26 nucleotides added to it
                                                 This is a c-myb mRNA target sequence (nucleotide no. 1919) of an enzymatic RNA molecule (ribozyme) which cleaves mRNA associated with the development or maintenance of a restenotic condition. The concr. of the ribozyme necessary to effect a therapeutic treatment is lower than that of an antisense oligonucleotide and the specificity of action is higher. (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                           Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           corresponding to 17 RNA polymerase promoter sequence. The resulting primer is described in AAT42249
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Detection of nucleic acid sequences - by polymerase chain reaction amplification, transcription using RNA polymerase and detection of RNA:DNA hybrids using antibodies.
                                                                                                                                                                                                                                                           0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Detection, PCR, polymerase chain reaction, hybrid; antibody; immunochemical detection; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match 0.8%; Score 14.2; DB 1; Length 21; Best Local Similarity 84.2%; Pred. No. 9.1e+02; Matches 16; Conservative 0; Mismatches 3; Indels
                                                                                                                                                                                                                     Score 14.2; DB 1; Length 21; Pred. No. 9.1e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Primer derived from hlyA gene used in modified PCR method.
                                                                                                                                                                                                                                                           Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 21 BP; 8 A; 1 C; 7 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                    Sequence 21 BP; 4 A; 5 C; 7 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                             0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example 1; Page 16; 31pp; English
                                                                                                                                                                                                                                                                                                859 GACCTGAAGCAGTACCTGG 877
                                                                                                                                                                                                                                                                                                                                   1 GCCTTGTAGCAGTACCTGG 19
               Claim 3; Page 20; 65pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                               AAT42247 standard; DNA; 21 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         94CA-02139070.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               94CA-02139070
                                                                                                                                                                                                                       0.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (first entry)
                                                                                                                                                                                                                                                               16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 1996-413110/42.
                                                                                                                                                                                                                       Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (BLAI/) BLAIS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               23-DEC-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          20-FEB-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CA2139070-A
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Blais BW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAT42247;
                                                                                                                                                                                                                                                                                                                                                                                            RESULT 1164
                                                                                                                                                                                                                                                               Matches
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This invention relates to novel oligonucleotide PCR primers which may be used to identify markers associated with genetic diseases and traits in dogs, in particular to disgnose genetic diseases that are not phenotypically visible and to identify carriers of recessive diseases. A specific application is diagnosis of copper toxicosis (CT). The invention can also be used to create a genetic map of the canine genome; to generate breed specific profiles; to establish paternity and to identify dogs from DNA fingerprints. The method provides rapid analysis of the target sequences from only a small sample of DNA. Diagnosis can be done at any time in the dog's life. The present sequence is that of a PCR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New oligonucleotide primers for diagnosis of genetic diseases and trait in dogs - amplify specific regions of the genome containing microsatellite repeats, especially for diagnosing copper toxicosis and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
                                                                                                                                                                                                                            genetic disease; genetic trait; dog; carrier of recessive disease; copper toxicosis, CT; canine genome map; breed-specific profile; DNA fingerprint; dog identification; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    .
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0.8%; Score 14.2; DB 1; Length 21; 34.2%; Pred. No. 9.1e+02; ve 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 21 BP; 7 A; 3 C; 6 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                Canine disease marker-related PCR primer 506
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Venta PJ, Yuzbasiyan-Gurkan V;
1503 TTCCATATTTGCACTAAAG 1521
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  855
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 1; Page 16; 40pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              N
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             BP.
                                                                                                          ВР
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TGTCTTTGAGTACCTGGAC
                            19 TICCATCTITCCACTAATG
                                                                                                                                                                                                                                                                                                                                                                                      97WO-US002396
                                                                                                                                                                                                                                                                                                                                                                                                                     96US-0012060P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    84.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                              (UNMI ) UNIV MICHIGAN. (UNMS ) UNIV MICHIGAN STATE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           21
                                                                                                          ADG77662 standard; DNA; 21
                                                                                                                                                                     (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAV51809 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 1997-435082/40.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Local Similarity
nes 16; Conserv
                                                                                                                                                                                                                                                                                           Canis familiaris.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        primer of the
                                                                                                                                                                    11-MAR-2004
                                                                                                                                                                                                                                                                                                                                                                                     18-FEB-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                     22-FEB-1996;
                                                                                                                                                                                                                                                                                                                           WO9731011-A1
                                                                                                                                                                                                                                                                                                                                                         28-AUG-1997.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Brewer GJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  837
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          carriers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAV51809
                                                                                                                                        ADG77662;
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Matches
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ID AAV5
XX
AC AAV5
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Gaps

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Murigneux A;

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Brassica species allele-specific oligonucleotide probes and primers useful for plant breeding.
                                                                              Sapolsky RJ,
                                                                                                                                                                   Example 1; Page 51; 65pp; English.
                                                                                                                                                                                                                                                                                                                             0.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Hudson T;
               96US-0032069P.
97US-00813507.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            97WO-US020313
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      96US-0030455P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAX09125 standard; DNA; 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first entry)
                                                                                                                                                                                                                                                                                                                                                          16; Conservative
                                                     (AFFY-) AFFYMETRIX INC
                                                                              Landry BS,
                                                                                                      WPI; 1998-333252/29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Wang D,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 1998-286974/25
                                                                                                                                                                                                                                                                                                                                             Best Local Similarity
               02-DEC-1996;
07-MAR-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   06-NOV-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         24-MAR-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            15-NOV-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   14-MAY-1998.
                                                                              Lemieux B,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Lander ES,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                         N
                                                                                                                                                                                                                                                                                                                                Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                               RESULT 1168
                                                                                                                                                                                                                                                                                                                                                         Matches
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                                                                                                                                                                                                                                                                                                                                                                                    AAV51705-V5200B are reverse PCR primers used to amplify fragments of the Zea mays genome in order to detect polymorphic markers. Such markers can be used in the construction of allele-specific primers and probes for amplification or hybridisation, e.g. to determine common or disparate ancestry between 2 or more plants, to monitor the genetic contribution of an ancestral plant, to trace the progeny of proprietary plants, in certification of a hybrid plant or to identify the progeny of a back-crossed plant with an ancestral plant
                                               Polymorphic marker, allele-specific; probe; amplification; PCR primer; hybridisation; plant; hybrid certification; genetic contribution; progeny; back-cross; hybrid; ancestry; corn; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Polymorphic marker; allele-specific; probe; amplification; PCR primer; hybridisation; plant; hybrid certification; genetic contribution; progeny; back-cross; hybrid; ancestry; corn; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
                                                                                                                                                                                                                                                                                                                            Brassica species allele-specific oligonucleotide probes and primers useful for plant breeding.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0.8%; Score 14.2; DB 1; Length 21;
84.2%; Pred. No. 9.1e+02;
tive 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                            Sapolsky RJ, Murigneux A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 21 BP; 7 A; 3 C; 5 G; 6 T; 0 U; 0 Other;
                        Zea mays genome reverse PCR primer #105.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Zea mays genome reverse PCR primer #108.
                                                                                                                                                                                                                                                                                                                                                                Example 1; Page 51; 65pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                587 CTGAGATTGGCTTTGGGAA 605
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CTGAGATTGGATTTGAAAA 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ВЪ
                                                                                                                                                                                       97WO-US021782
                                                                                                                                                                                                              96US-0032069P.
97US-00813507.
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02-FEB-1999 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAV51812 standard; DNA; 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                                                                                                                                                                                                            Landry BS,
                                                                                                                                                                                                                                                   (AFFY-) AFFYMETRIX INC
                                                                                                                                                                                                                                                                                                   WPI; 1998-333252/29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Best Local Similarity
                                                                                                                                                                                                             02-DEC-1996;
07-MAR-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                01-DEC-1997;
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                                                                                                                                    WO9824796-A1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         16;
                                                                                                                                                              11-JUN-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      02-FEB-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         11-JUN-1998
                                                                                                Synthetic.
                                                                                                                                                                                                                                                                            Lemieux B,
                                                                                                               Zea mays.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
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AAV51705-V52008 are reverse PCR primers used to amplify fragments of the Zea mays genome in order to detect polymorphic markers. Such markers can be used in the construction of allele-specific primers and probes for amplification or hybridisation, e.g. to determine common or disparate ancestry between 2 or more plants, to monitor the genetic contribution of an ancestral plant, to trace the progeny of proprietary plants, in certification of a hybrid plant or to identify the progeny of a backcrossed plant with an ancestral plant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Polymorphism; biallelic; human; forensic; paternity testing; disease; detection; phenotypic typing; characteristic; infection; hereditary; autoimmune disease; cancer; inflammation; drug; therapy; medicament; treatment; marker; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         .
0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 14.2; DB 1; Length Pred. No. 9.1e+02; 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human biallelic polymorphic marker upstream primer #5.
                                                                                                                                                                                                                                                                                                                                                                  Sequence 21 BP; 7 A; 3 C; 5 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           testing or phenotypic typing for disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              RES.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 15; Page 46; 310pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          587 CTGAGATTGGCTTTGGGAA 605
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AAX09121-X10268 are allele-specific oligonucleotide primers used in the isolation of various biallelic polymorphic markers found in the human egenome (represented in AAX10269-X12937). These primers can be used in a method for determining polymorphic forms in an individual for use in e.g. forensics, paternity testing or for phenotypic typing for diseases such as agammaglobulinemia, diabetes insipidus, Lesch-Nyham syndrome, muscular cyptrophy, Wiskott-Aldrich syndrome, Pabry's disease, familial cyptrophy, Wiskott-Aldrich syndrome, Pabry's disease, hereditary hypercholesterolemia, polycystic Kidney disease, hereditary cyptropis, von Willebrand's disease, tubercous sclerosis, hereditary haemorrhagic telangiectasia, familial colonic polyposis, Ehlers-Danlos syndrome, osteogenesis imperfecta, acute intermittent porphyria, autoimmume diseases, inflammation, cancer, diseases of the nervous system, infection by pathogenic microorganisms, and characteristics such as longevity, appearance (e.g. baldness, obseity), strength, speed, endurance, ferrility, and susceptibility or receptivity to particular crugs or therapeutic treatments. The isolated polymorphic nucleic acid segments can also be used to produce medicaments for the treatment or prophylaxis of such diseases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ATP binding cassette, ABC transporter, ABCR; Stargardt Disease, therapy; Fundus Flavimaculatus; age-related macular degeneration; diagnosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Li Y;
Smallwood PM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Retina-specific ATP-binding cassette transporter and DNA - useful for, e.g. diagnosis and treatment of macular degeneration, such as in Stargardt Disease, Fundus Flavimaculatus and age-related degeneration.
                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
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Singh N,
                                                                                                                                                                                                                                                                                                                                                                                       Score 14.2; DB 1; Length 21; Pred. No. 9.18+02; 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PCR primer ABCR.EXON31:F for ABCR coding sequence.
                                                                                                                                                                                                                                                                                                                                                        Sequence 21 BP; 7 A; 2 C; 10 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Leppart M,
Shroyer NF,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (UYJO ) UNIV JOHNS HOPKINS.
(USSH ) US DEPT HEALTH & HUMAN SERVICES.
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thans J, Rattner A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1678 CCCAACTACATCTTCCCTG 1696
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 41; Page 30; 79pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      crecachacererrecers 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAV08249 standard; DNA; 21 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              98WO-US003895.
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                                                                                                                                                                                                                                                                                                                                                                                          Ouery Match 0.8%;
Best Local Similarity 84.2%;
Matches 16; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 1998-495375/42.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PCR primer; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Homo sapiens.
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Lupski JR,
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(USSH)
(UTAH)
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This sequence represents a PCR primer for DNA encoding the human retina

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ANY62007-V62009 are PCR primers used in a novel method for the enhanced detection of DNA sequences, via a nucleic acid amplification procedure, especially for detecting pathogens. Minute samples of pathogens (c. 10 cells) cannot be detected effectively by PCR. The minute quantities of product formed by PCR are then transcribed into RNA enhancement products, which further amplifies the target sequences to detectable levels. Detection then takes place with antibodies for DNA:RNA hybrids, which enable detection if the product volume formed is still small, but is specific enough just for this type of product. The method is especially useful for detecting the following pathogens: Listeria monocytogenes, L. innocua, L. ivanovii, L. seeligeri, L. welshimeri, L. murrayi, L. grayi, inconcus thermophillus, Lactobacillus casel, Lactococcus lactis, Micrococcus thermophillus, Lactobacillus casel, Lactococcus epidermidis, Micrococcus Luteus, Enterococcus epidermidis, Bacillus subtilis, Pseudomonas aeruginosa, Escherichia acil, Salmonella typhimurium, or Yersinia enterocolitica. (Updated on 25-
                                                                                                                                                                                            ô
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Detection; pathogen; amplification; RNA enhancement product; PCR primer; DNA/RNA hybrid; Listeria sp; Streptococcus sp; Lactobacillus sp; Lactococcus sp; Micrococcus sp; Entercoccus sp; Stapylococcus sp; Bacillus sp; Pseudomonas sp; Escherichia coli; Salmonella typhimurium; Yersinia enterocolitica; ss.
              may be used in compositions for screening agents that alters ABCR. The agent can inhibit Stargardt Disease, Fundus Flavimaculatus and agerelated macular degeneration (MD). Primers (such as this sequence) and probes for the ABCR DNA can be used in a diagnostic kit for detecting MD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Enhanced detection by nucleic acid amplification, especially of Listeria - uses formation of DNA-RNA hybrids after amplification, and then
                                                                                                                                                                                          Gaps
specific ATP binding cassette transporter (ABCR) of the invention.
                                                                                                                                                                                            ·.
                                                                                                                                                      Length 21;
                                                                                                                                                                                            3; Indels
                                                                                                                BP; 6 A; 8 C; 3 G; 4 T; 0 U; 0 Other;
                                                                                                                                                      Score 14.2; DB 1;
Pred. No. 9.1e+02;
                                                                                                                                                                                          0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         L monocytogenes hlyA gene PCR primer A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     specific immuno-detection of these.
                                                                                                                                                                                                                                  1389 CCTCACCAAGCTGTTGCAG 1407
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Example 1; Col 12; 15pp; English.
                                                                                                                                                                                                                                                                       3 carcaccadererrecad 21
                                                                                                                                                                                                                                                                                                                                                                       BP.
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94US-00366619.
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                                                                                                                                                          0.8%;
                                                                                                                                                                        1 Similarity 84.2%;
16; Conservative
                                                                                                                                                                                                                                                                                                                                                                       AAV62007 standard; DNA; 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Synthetic.
Listeria monocytogenes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                   (revised)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 1998-593985/50.
                                                                                                                                                                            Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (KALY-) KALYX
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  23-SEP-1996;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           30-DEC-1994;
                                                                                                                    Sequence 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                 25-MAR-2003
11-JAN-1999
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                                                                                                                                                                                                                                                                                                                                                                                                             AAV62007;
                                                                                                                                                          Query Match
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Gaps

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This invention describes a novel method for identifying an inhibitor of potentially useful for treatment of cancer, where the inhibitor is active on a gene vital for cell growth or viability, and where the gene is subject to loss of heterozygosity (LOH) in a cancer. The inhibitor is used for preventing the development of cancer in a patient having a precancerous condition, by administering to the patient a first allele specific inhibitor (ASI) targeted to an allele of a first essential gene present in cells of the precancerous condition, where the normal somatic cells of the precancerozygous for the first gene, the inhibitor is active on at least one but less than all allelic forms of the gene present in a population and targets only one allelic form present in the cormal somatic cells, and the first gene. The products and methods can be used in the diagnosis, prevention and treatment of LOH disorders, e.g. cancers, atherosclerotic plaques, premalignant metaplastic or dysplastic essions, benign tumours, endometriosis, polycystic kidney disease, and graft versus host disease. The method can also be used to remove malignant cells from bone marrow transplants. AAZSSB12-ZSG825 represent human polymorphic sites described in the method of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Polymorphism; human; inhibitor; cancer; treatment; cell growth; LOH; cell viability; loss of heterozygosity; precancerous condition; ASI; allele specific inhibitor; somatic cell; diagnosis; prevention; atherosclerotic plaque; premalignant metaplastic lesion; endometriosis; dysplastic lesion; benign tumour; polycystic kidney disease; transplant; graft versus host disease; malignant cell removal; bone marrow; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Identifying target genes for allele-specific drugs - used for diagnosis, prevention and treatment of, e.g. cancers, atherosclerotic plaque, dysplastic lesions, endometriosis or graft versus host disease.
                                                                                                                                   0;
                                                                                   0.8%; Score 14.2; DB 1; Length 21;
84.2%; Pred. No. 9.1e+02;
iive 0; Mismatches 3; Indels
                                        Sequence 21 BP; 8 A; 1 C; 7 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Ledley FD, Stanton VP;
                                                                                                                                                                          1503 TICCATATITGCACTAAAG 1521
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MAR-2003 to correct PR field.)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human polymorphic region 313
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     97US-0041057P
                                                                                                                                                                                                                                                                                                                               AAZ26124 standard; DNA; 21
                                                                                                                                                                                                                                                                                                                                                                                                                      30-NOV-1999 (first entry)
                                                                               Ouery Match
Best Local Similarity 84.2
Matches 16; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 1998-521232/44.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                           AAZ26124;
                                                                                                                                                                                                                                                                                    RESULT 1171
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                                                                                                                                                                                                                                                                                                                                                                                                                           cell viability; loss of heterozygosity; precancerous condition; ASI; allele specific inhibitor; somatic cell; diagnosis; prevention; atherosclerotic plaque; premalignant metaplastic lesion; endometriosis; dysplastic lesion; benign tumour; polycystic kidney disease; transplant; graft versus host disease; malignant cell removal; bone marrow, ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Identifying target genes for allele-specific drugs - used for diagnosis, prevention and treatment of, e.g. cancers, atherosclerotic plaque, dysplastic lesions, endometriosis or graft versus host disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     malignant cells from bone marrow transplants. AA225812-226825 represent human polymorphic sites described in the method of the invention
                                              Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             This invention describes a novel method for identifying an inhibitor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   graft versus host disease. The method can also be used to remove
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    Length 21;
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                                            Indels
                                                                                                                                                                                                                                                                                                                                                                                                        Polymorphism; human; inhibitor; cancer; treatment;
  Score 14.2; DB 1; Pred. No. 9.1e+02; 0; Mismatches 3;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Stanton VP;
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                                                                                 940 GGCCTGGCCTACTGCCACC 958
                                                                                                                           GCCCTGGCCTTCCGCCACC 21
                                                                                                                                                                                                                                     BP
                                                                                                                                                                                                                                                                                                                                                              Human polymorphic region 431.
  0.8%;
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                                                                                                                                                                                                                                  AAZ26242 standard; DNA; 21
                                                                                                                                                                                                                                                                                                                  30-NOV-1999 (first entry)
Query Match 0.8
Best Local Similarity 84.2
Matches 16; Conservative
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                                                                                                                                                                                                                                                                            AAZ26242;
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                                                                                                                                                                                                                 AAZ26242/c
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Gaps

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3; Indels

0; Mismatches

Best Local Similarity 84.2%; Matches 16; Conservative

Sequence 21 BP; 2 A; 12 C; 4 G; 3 T; 0 U; 0 Other;

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Antisense; oligonucleotide; immediate early; DNA polymerase; C
cytomegalovirus; inhibition; replication; sugar modification;
phosphorothioate; infection; retinitis; ss.
                                                                                                                             Anti-CMV oligonucleotide #2922
                                      AAX17882 standard; DNA; 21
                                                                                                11-MAY-1999 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (ISIS-) ISIS PHARM INC.
                                                                                                                                                                                                                                  Human herpesvirus 5.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 1998-568330/48.
                                                                                                                                                                                                                                                                 Key
modified_base
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                                                                                                                                                                                                                                                                                                                                                                                    15-0CT-1998.
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                                                                                                                                                                                                                      Synthetic
                                                                     AAX17882;
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         RESULT 1174
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                           AAX17882
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      potentially useful for treatment of cancer, where the inhibitor is active on a gene vital for cell growth or viability, and where the gene is subject to loss of heterozygosity (LOH) in a cancer. The inhibitor is precancerous condition, by administering to the patient a first allele specific inhibitor (ASI) targeted to an allele of a first essential gene present in cells of the precancerous condition, where the normal somatic cells of the patient are heterozygous for the first gene, the inhibitor is active on at least one but less than all allelic forms of the gene present in a population and targets only one allelic forms of the gene hormal somatic cells, and the first gene. The products and methods can be used in the diagnosis, prevention and treatment of LOH disorders, e.g. cancers, atherosclerotic plaques, premalignant metaplastic or dysplastic lessions, benign tunours, endometriosis, polycystic kidney disease, and
                                                                                                                                                                                                                   Polymorphism; human; inhibitor; cancer; treatment; cell growth; LOH; cell viability; loss of heterozygosity; precancerous condition; ASI; allele specific inhibitor; somatic cell; diagnosis; prevention; atherosclerotic plaque; premalignant metaplastic lesion; endometriosis; dysplastic lesion; benign tumour; polycystic kidney disease; transplant; graft versus host disease; malignant cell removal; bone marrow; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Identifying target genes for allele-specific drugs - used for diagnosis, prevention and treatment of, e.g. cancers, atherosclerotic plaque, dysplastic lesions, endometriosis or graft versus host disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    graft versus host disease. The method can also be used to remove malignant cells from bone marrow transplants. AAZ25812-Z26825 represent human polymorphic sites described in the method of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           invention describes a novel method for identifying an inhibitor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0.8%; Score 14.2; DB 1; Length 21;
84.2%; Pred. No. 9.1e+02;
iive 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Seguence 21 BP; 1 A; 5 C; 10 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Stanton VP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Disclosure; Fig 7; 605pp; English.
235 GGTGGTGGCGGCAGTGACC 253
                             N
                                                                                                      BP.
                                                                                                                                                                                                                                                                                                                                                                                                                        98WO-US005419.
                    GGTGGTGGCGCCACTCAGC
                                                                                                                                                                                           Human polymorphic region 291.
                                                                                                                                                                                                                                                                                                                                                                                                                                                  97US-0041057P
                                                                                                        AAZ26102 standard; DNA; 21
                                                                                                                                                               (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (VARI-) VARIAGENICS INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Ledley FD,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 1998-521232/44
                                                                                                                                                                                                                                                                                                                                                                                                                        19-MAR-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                  20-MAR-1997;
                                                                                                                                                                                                                                                                                                                                 Homo sapiens
                                                                                                                                                                                                                                                                                                                                                             WO9841648-A2
                                                                                                                                                                 30-NOV-1999
                                                                                                                                                                                                                                                                                                                                                                                           24-SEP-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Housman D,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
                             20
                                                                                                                                    AAZ26102
                                                                            1173
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Chapman S;

Anderson KP,

Kisner DL,

98WO-US006895. 97US-00838715

/*tag= a /note= "contains phosphorothioate internucleotide

linkages"

Location/Qualifiers 1. .21

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0
                                                                                                                                                                                                                                               replication. Optionally the oligonucleotides include at least one 2'-(2-methoxy) sugar modification or phosphorothioate internucleotide linkages. The oligonucleotides are used to inhibit CMV infections (by in vivo or in vitro contact with cells, tissues or body fluids), especially
New antieense oligonucleotides that target cytomegalovirus nucleic acid particularly including 2-methoxyethoxy sugar modifications, especially for treating viral retinitis, with long-lasting retention in the retina.
                                                                                                                                                            Antisense oligonucleotides (AAX17861-X17924) are targeted to a nucleic acid (AAX17925-X17948) encoding IE (immediate early) 1 or 2, or DNA polymerase of cytomegalovirus (CMV) and are able to inhibit CMV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human integrin beta 3 quantitative real-time PCR primer, SEQ ID NO:3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 21;
                                                                                                                                                                                                                                                                                                                                                                   to treat or prevent CMV infections, particularly retinitis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 14.2; DB 1; Length 2
Pred. No. 9.1e+02;
0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 21 BP; 0 A; 7 C; 4 G; 10 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            131 GGATGAAGAAGATCAAACG 149
                                                                                                                   Claim 2; Page 24; 99pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                20 GCAAGAAGAGAGCAAACG 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0.8%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match 0.8
Best Local Similarity 84.2
Matches 16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     03-JUL-2000
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Gaps

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217 GGCCTGGATGAGAGTGGTG 235

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Best Local Similarity Matches 16; Conserv

Single nucleotide polymorphism; SNP; STP2; phenol sulphotransferase; probe; genotyping; human; drug metabolism; ss.

Homo sapiens

Human STP2 gene promoter polymorphism sequence 108.

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Integrin beta 3; human endothelial glycoprotein; GP3A; GPIIIa; ITGB3; CD61; platelet glycoprotein 3a; cellular adhesion; vitronectin receptor; fibronectin receptor; expression inhibition; antisense therapy; tumour formation; cancer invasion; bleeding disorder; inflammation; quantitative real-time PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 21 BP; 5 A; 5 C; 6 G; 5 T; 0 U; 0 Other;
                                                                                                                                                           Monia BP;
                                                                                                                                                                                                                    Example 13; Col 39; 33pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  614 CCTACATTAAGCTGGACAA 632
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1 CCGTCATTAGGCTGGACAA 19
                                                                                                          99US-00344520.
                                                                                                                          99US-00344520,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAZ59350 standard; DNA; 21
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                                                                                                                                                          Bennett CF, Cowsert LM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                                          (ISIS-) ISIS PHARM INC
                                                                                                                                                                           WPI; 2000-246189/21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Local Similarity
nes 16; Conserv
                                                          Homo sapiens
                                                                                                         25-JUN-1999;
                                                                                                                         25-JUN-1999;
                                                                         US6037176-A.
                                                                                          14-MAR-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
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Matches
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Sequences AAZ59305-Z59352 are fragments of the human STP2 gene. The fragments are from the 8 exons, the promoter region, 3' and 5' untranslated regions of the STP2 gene. Each sequence contains a newly identified STP2 gene single nucleotide polymorphism (SNP). STP2 is a phenol sulphotransferase. Substrates for STP2 include minoxidil, actenaninophen, and paranitrophenol. Several of the modified changes identified at the polymorphism sites, give rise to an amino acid change may result in altered enzyme activity. The sequences of mino acid changes for detecting STP2 polymorphisms. The polymorphic probes are used in screening and genotyping, i.e. to predict the rate of metabolism of STP2 substrates, potential drug-drug interactions and adverse side effects. They can also be used to detect diseases resulting from accidental or occupational expense to toxins and to establish animal, cell or in vitro models for drug metabolism
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human biallelic marker downstream amplification primer SEQ ID NO:8100.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Novel nucleic acid used for genotyping, e.g. to predict rate of drug
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human genome; biallelic marker, high density disequilibrium map; genomic map; haplotype; polymorphic base; genotyping; haplotyping; hybridisation; identification; characterisation; amplification; single nucleotide polymorphism; SNP; PCR primer; amplification; single nucleotide polymorphism; SNP; PCR primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 14.2; DB 1; Length 21;
Pred. No. 9.1e+02;
0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 21 BP; 2 A; 9 C; 2 G; 8 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                    /*tag= a
/note= "Site of polymorphism"
                                                                                                                                                                                                  Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 2; Page 17; 46pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     44
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local Similarity 84....
Local 16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAZ73744 standard; DNA; 21
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (AXYS-) AXYS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2000-105892/09.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Guida M, Kurth J;
                                                                                                                                                                                                                                                                                                                                                                                                                            199-JUN-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              10-JUN-1998;
                                                                                                                                                                                                                                                                                                                              WO9964630-A1
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                                                                                                                                                                                                                                                                                                                                                                          16-DEC-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                metabolism.
                                                                                                                                                                                                                          variation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAZ73744;
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g
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequences AAA07029-A07030 represent human integrin beta 3 PCR primers used in quantitative real-time PCR with probe AAA07031 in an exemplification of the present invention. The invention relates to antisense oligonucleotides targetted to the human integrin beta 3 gene, which inhibit its expression. A series of oligonucleotides (AAA07035-AAA07074) were designed to target different regions of the human integrin beta 3 RNA, and were analysed for their effect on integrin beta 3 mRNA, levels by quantitative real-time PCR. GAPBH (qlyceraldehyde-3-phosphate) mRNA levels were measured as a control. Integrins constitute one of four Classes of cellular adhesion molecules, and play an important role in cell migration, cell anchorage to substrates and cytoadhesion signalling pathways. They are heterodimeric cation-dependent membrane glycoproteins composed of an alpha and beta subunit. Integrin beta 3 (also known as human endothelial glycoprotein, GP3A, GPIIIa, ITGB3, CD61 and platelet glycoprotein 3a) is the common beta subunit partner of the members of the beta-3 subfamily of integrins. This family consists of the vitronectin ceptor (alpha-V-beta-3) and the fibronectin receptor (alpha-IIb-beta-23). Cells expressing this class of integrin can adhere to various matrix process. Integrin beta 3 is implicated in conditions such as vascular response.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          restenosis, excessive bone resorption, angiogenesis (in melanoma), tumour invasion, platelet aggregation and Glanzmann's thrombasthenia. The oligomucleotides of the invention are useful for diagnosis, prevention and treatment of conditions associated with integrin beta 3 expression, such as tumour formation, inflammation, infections and the diseases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New antisense compound that inhibits human integrin beta3, useful e.g. for treating or preventing infection, inflammation and tumors.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
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84.2%; Pred. No. 9.1e+02;
tive 0; Mismatches 3; Indels
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Gaps

0;

diagnosis; ss

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10-SEP-1999; 99US-0153357P.
26-JUL-2000; 2000US-0220947P.
89US-00399728
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                                                                                                          (PALE/) PALESE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens
              21-NOV-1989;
22-MAY-1990;
                                            04-AUG-1992;
01-FEB-1994;
 28-AUG-1989;
                                                                            01-JUN-1994;
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                                                                                                                                                      Palese
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              RESULT 1179
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                                                                                                                                                                                                                                                                                                                                                  AAZ65654 to AAZ69578 represent human biallelic markers from the present invention, which contain a polymorphic base at position 24 of their nuclectide sequences. AAZ69579 to AAZ7740 represent amplification primers for the biallelic markers. The biallelic markers of the invention have a variety of uses: they can be used for high density mapping of the mann genome, and in complex association studies and haplotyping studies which are useful in determining the genetic basis for disease states. Compositions and methods of the invention can also be useful for the identification of the targets for the development of pharmaceutical agents and diagnostic methods, as well as the characterisation of the pharmaceutical agents acting on a disease as well as other treatment. W.B. The SEQ ID NOS 2852, 2913, 2974, 3035, 3096, 3157, 3227, 3297 and 3367, are not actually given a sequence in the Sequence Listing from the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Recombinant negative strand viral RNA template, virus particle, RNA directed RNA polymerase complex, expression, chimeric virus; vaccine;
                                                                                                                                                                                                                                                                            Novel biallelic markers used to construct a high density disequilibrium
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0.8%; Score 14.2; DB 1; Length 21;
84.2%; Pred. No. 9.1e+02;
ative 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 21 BP; 7 A; 4 C; 5 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                    Chumakov I;
                                                                                                                                                                                                                                                                                                                          Claim 8; Page 1957; 2745pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               392 CGGATGAGGTGCAGTCTCC 410
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       21 CAGATGATTTGCAGTCTCC 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAZS6234 standard; DNA; 21 BP.
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                                                                                                             99WO-IB000822
                                                                                                                                           98US-0082614P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 16; Conservative
                                                                                                                                                                                                                      Blumenfeld M,
                                                                                                                                                                                                                                                                                             map of the human genome
                                                                                                                                                                                                                                                   WPI; 2000-013267/01
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Influenza virus.
Synthetic.
                                                                                                                                                                                     (GEST ) GENSET
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                                                                                                                                                          23-NOV-1998;
                    Homo sapiens
                                                 WO9954500-A2
                                                                                                             21-APR-1999;
                                                                                                                                           21-APR-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          packaging;
                                                                                                                                                                                                                    Cohen D,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
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The present invention describes a chimeric virus comprising influenza virus containing a heterologous RNA segment from another strain of influenza virus or 8 genomic segments from different strains of influenza virus, with each segment comprising the reverse complement of a mRNA coding sequence operatively linked to a binding site specific for an RNA-directed RNA polymerase of a negative strand RNA virus. The recombinant negative strand virus RNA templates may be used to express heterologous gene products in appropriate host cell systems and/or to construct recombinant viruses that express, package and/or present the heterologous gene product. The expression products and chimeric viruses may be used in vaccine formulations. ANYS7746 to AAX556234 to AAX56290, represent sequences used in the exemplification of the present invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                  Chimeric virus containing influenza virus RNA segments, useful for expressing heterologous gene products in appropriate host cell systems.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
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/standard_name= "single nucleotide polymorphism"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 21 BP; 6 A; 3 C; 5 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human gene single nucleotide polymorphism #2298.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                908 ACGTGAAACTGTTCCTGTT 926
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2 ACGAGGAAATGTTCCTGTT 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example; Col 55; 67pp; English.
89US-00440053.
90US-00527237.
92US-00925061.
94US-00190698.
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                                                                                                                                    94US-00252508
                                                                                                                                                                                                                                                                                                            Garcia-Sastre A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAF97537 standard; DNA; 21
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Best Local Similarity 84.2'
                                                                                                                                                                                                                                      (GARC/) GARCIA-SASTRE
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WPI; 2001-226749/23
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 10-SEP-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    15-MAR-2001.
            Lander ES,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Lander ES,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Variation
                                                                                                                                                                                                                                                                                                                                                                                                                                           AAF96385;
                                                                                                                                                                                                                                                                                              Query Match
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                                                                                                                                                                                                                                                                                                                                                            g
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                                                                                                                                                                                                             genes. The sequences at a number of polymorphic sites are also provided in the specification. In particular, the method can be used in the diagnosis of atherosclerosis, mycoradial infarction, coronary heart disease, stroke, perhipheral vascular diseases, venous thromboembolism and pulmonary embolism. Single nucleotide polymorphisms (SNPs) are also correlations to diseases. The present sequence is an example of one of the human gene SNPS shown in the specification
                                                                                                                                                                            The present invention provides a method of diagnosing a vascular disease in an individual, involving determining the sequence at various polymorphic sites within the human thrombospondin 1 and thrombospondin 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human; variant thrombospondin 1; variant thrombospondin 4; SNP; polymorphism; vascular disease; coronary artery disease; forensics; myocardial infarction; atherosclerosis; stroke; venous thromboembolism; pulmonary embolism; paternity test; ds.
                                                        Mccarthy JJ;
                                                                                                 Nucleic acids comprising single nucleotide polymorphisms, useful in applications such as forensics, paternity testing, medicine, genetic analysis and phenotype correlations to diseases such as diabetes and
                                                                                                                                                                                                                                                                                                                                                              Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /standard_name= "single nucleotide polymorphism"
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0
                                                                                                                                                                                                                                                                                                                                      0.8%; Score 14.2; DB 1; Length 21; llarity 84.2%; Pred. No. 9.1e+02; Conservative 0; Mismatches 3; Indels
                                                        Daley GQ,
                                                                                                                                                                                                                                                                                                                  Sequence 21 BP; 3 A; 6 C; 9 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human gene single nucleotide polymorphism #73.
                                                      Bolk S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (WHED ) WHITEHEAD INST BIOMEDICAL RES. (MILL-) MILLENNIUM PHARM INC.
                     (WHED ) WHITEHEAD INST BIOMEDICAL RES. (MILL-) MILLENNIUM PHARM INC.
                                                      Gargill M, Ireland JS,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Location/Qualifiers
                                                                                                                                                          Example, Page 204; 242pp; English
                                                                                                                                                                                                                                                                                                                                                                                   490 GACATCCGGCTGCCTGAGG 508
                                                                                                                                                                                                                                                                                                                                                                                               21 GCCTCCGCTGCCTGAGG 3
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26-JUL-2000; 2000US-0220947P.
16-AUG-2000; 2000US-0225724P.
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16-AUG-2000; 2000US-0225724P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAF95312 standard; DNA; 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
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                                                                           WPI; 2001-226749/23
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nes 16; Conserv
                                                                                                                                    atherosclerosis.
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                                                      ES,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Key
Variation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAF95312;
                                                                                                                                                                                                                                                                                                                                       Query Match
                                                      Lander
                                                                                                                                                                                                                                                                                                                                                                                                                                          RESULT 1180
                                                                                                                                                                                                                                                                                                                                                           Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human; variant thrombospondin 1; variant thrombospondin 4; SNP; polymorphism; vascular disease; coronary artery disease; forensics; myocardial infarction; atherosclerosis; stroke; venous thromboembolism; pulmonary embolism; paternity test; ds.
                                                                                                        Nucleic acids comprising single nucleotide polymorphisms, useful in applications such as forensics, paternity testing, medicine, genetic analysis and phenotype correlations to diseases such as diabetes and atherosclerosis.
   Mccarthy JJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /standard_name= "single nucleotide polymorphism"
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Pred. No. 9.1e+02;
0; Mismatches 3; Indels
Daley GQ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Daley GQ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Seguence 21 BP; 9 A; 5 C; 4 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          duman gene single nucleotide polymorphism #1146.
   Bolk S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Bolk S,
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   Ireland JS,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Location/Qualifiers replace (11, A)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Ireland JS,
                                                                                                                                                                                                                                                                         Example; Page 51; 242pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        719 AACATGAAGAGGGGCACC 737
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAF96385 standard; DNA; 21 BP
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2000US-0220947P.
2000US-0225724P.
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   Gargill M,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Best Local Similarity
                                                            WPI; 2001-226749/23
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Gaps

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Length 21;

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contain single nucleotide polymorphisms (SNPs). A method is included in the invention for analysing a nucleic acid sample, which consists of determining the base occupying any one of the polymorphic sites given in the SNP containing sequences. The nucleotide sequences can be used in the diseases, diseases of the carditovascular system, and infection by microorganisms. The oligonucleotides are also useful in the manufacture of a medicament for the treatment or prophylaxis of the diseases, and as a pharmacutical. SNP containing oligonucleotides are also useful in an asphications such as phenotype correlation, forensics, paternity testing, medicine and genetic analysis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DNA sequences AAH62100 - AAH62688 represent segments of human genes which contain single nucleotide polymorphisms (SNPs). A method is included in the invention for analysing a nucleic acid sample, which consists of determining the base occupying any one of the polymorphic sites given in the SNP containing sequences. The nucleotide sequences can be used in the
sequences AAH62100 - AAH62688 represent segments of human genes which
                                                                                                                                                                                                                   Revised record issued on 09-SEP-2004 : Correction to Feature Table Key
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Single nucleotide polymorphism; SNP; human; cancer; inflammation; heart disease; paternity testing; forensic science; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New nucleic acid segments of the human genome, particularly from including polymorphic sites, for phenotype correlation, forensics, paternity testing, medicine and genetic analysis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Opiate receptor like 1 polymorphism containing DNA fragment #538.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /*tag= a
/standard name= "single nucleotide polymorphism"
                                                                                                                                                                                                                                                                                                                     3; Indels
                                                                                                                                                                                                                                                   Sequence 21 BP; 3 A; 4 C; 11 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                  Score 14.2; DB 1;
Pred. No. 9.1e+02;
                                                                                                                                                                                                                                                                                                                     0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                      39 GGCAGGAGGACCAGCAGTG 57
                                                                                                                                                                                                                                                                                                                                                                                      1 eccecadeccercedare 19
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                                                                                                                                                                                                                                                                                  0.8%;
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                                                                                                                                                                                                                                                                                                                         16; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (revised)
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                                                                                                                                                                                                                                                                                    Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                          RESULT 1183
                                                                                                                                                                                                                                                                                                                         Matches
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                                                                                                                                                                                                                                   disease, stroke, peripheral vascular diseases, venous thromboembolism and pulmonary embolism. Single nucleotide polymorphisms (SNPs) are also useful in forensics, paternity testing, genetic analysis and phenotype correlations to diseases. The present sequence is an example of one of the human gene SNPS shown in the specification
                                                                                                                                 The present invention provides a method of diagnosing a vascular disease in an individual, involving determining the sequence at various polymorphic sites within the human thrombospondin 1 and thrombospondin 4 genes. The sequences at a number of polymorphic sites are also provided in the specification. In particular, the method can be used in the diagnosis of atherosclerosis, myocardial infarction, coronary heart
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New nucleic acid segments of the human genome, particularly from genes including polymorphic sites, for phenotype correlation, forensics, paternity testing, medicine and genetic analysis
                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
               Nucleic acids comprising single nucleotide polymorphisms, useful in applications such as forensics, paternity testing, medicine, genetic analysis and phenotype correlations to diseases such as diabetes and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Single nucleotide polymorphism; SNP; human; cancer; inflammation; heart disease; paternity testing; forensic science; ds.
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/standard_name= "single nucleotide polymorphism"
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                                                                                                                                                                                                                                                                                                                                                                      0.8%; Score 14.2; DB 1; Length 21; 84.2%; Pred. No. 9.1e+02;
                                                                                                                                                                                                                                                                                                                                                                                                         Indels
                                                                                                                                                                                                                                                                                                                                       Sequence 21 BP; 5 A; 4 C; 7 G; 5 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                         0; Mismatches
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                                                                                                     Example; Page 130; 242pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                          1167 GGGCTGCATCTTCTATGAG 1185
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1 GGGCATCAGCTTCTATGAG 19
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (revised)
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                                                                                                                                                                                                                                                                                                                                                                                        Best Local Similarity
                                                                    atherosclerosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WO200138576-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Homo sapiens
Unidentified
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     24-NOV-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               09-SEP-2004
12-SEP-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    31-MAY-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Cargill M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Key
variation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAH62348;
                                                                                                                                                                                                                                                                                                                                                                          Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             RESULT 1182
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1195 GGCCGTCCCTCTTTCCGG

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   useful
diagnosis or monitoring of diseases, such as cancer, inflammation, heart diseases, diseases of the cardiovascular system, and infection by microorganisms. The oligonucleotides are also useful in the manufacture of a medicament for the treatment or prophylaxis of the diseases, and as a pharmaceutical. SNP containing oligonucleotides are useful in applications such as phenotype correlation, forensics, paternity testing,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mouse; ztrypl; serine protease; tryptase; inflammation; fertilisation; cardiovascular disease; infertility; asthma; immune disorder; stroke; gastrointestinal disorder; testicular function; contraceptive;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The present invention provides the protein and coding sequences of the
                                                                                                                                                                                                                                                                         Revised record issued on 09-SEP-2004 : Correction to Feature Table Key
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   mouse serine protease polypeptides ztrypl and polynucleotides, use treating cardiovascular disease, infertility, impotence and other
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             addition, they can be used to modulate testicular function and as
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0
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84.2%; Pred. No. 9.1e+02;
                                                                                                                                                                                                                                                                                                                                                                                                          Length 21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Murine ztrypl coding sequence PCR primer ZC18,365.
                                                                                                                                                                                                                                                                                                                                        Sequence 21 BP; 3 A; 6 C; 8 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 21 BP; 1 A; 9 C; 4 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                      0.8%; Score 14.2; DB 1;
34.2%; Pred. No. 9.1e+02;
ve 0; Mismatches 3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Example 1; Page 102; 112pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TGTTCCAGCTGCTCCGTGG 941
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                                                                                                                                                                                                          medicine and genetic analysis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TGATCCGGCGGCTCCGTGG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              09-AUG-2000; 2000WO-US022156.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 99US-00376445
                                                                                                                                                                                                                                                                                                                                                                                                                                         84.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAF75649 standard; DNA; 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                         Local Similarity 84.2
hes 16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (ZYMO ) ZYMOGENETICS INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Best Local Similarity 84.2
Matches 16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Presnell SR, Taft DW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2001-202859/20.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New mouse serine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WO200112788-A2.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PCR primer; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             18-AUG-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           musculus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  10-MAY-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          UDP-GAG is useful for obtaining transgenic cyanogenic plants with altered resistance to pathogens and herbivore responses
                                                                                                                                                                                            UDP-glucose:aglycon-glucosyltransferase; UDP-GAG; cyanohydrin; terpenoid;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DNA molecule coding for UDP-glucose:aglycon-glucosyltransferase conjugating cyanohydrin, terpenoid or phenylderivative to glucose, for producing transgenic plants having modified cyanogenic glucoside
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
                                                                                                                                                                 PCR primer for UDP-glucose:aglycon-glucosyltransferase DNA probe.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ö
                                                                                                                                                                                                         glucose, transgenic plant; cyanogenic glucoside biosynthesis; pathogen resistance, herbivore response; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0.8%; Score 14.2; DB 1; Length 21; 4.2%; Pred. No. 9.1e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human RecQ5 type DNA helicase sequencing primer 501.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 21 BP; 3 A; 5 C; 13 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                         UNIV ROYAL VETERINARY & AGRIC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 4; Page 17; 31pp; English.
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          20
                                                                                BP
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99EP-00123838
                                                                                                                                                                                                                                                                                                                                                                                                                                       Jones
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          84.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAF87687 standard; DNA; 21
                                                                                21
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            16; Conservative
                                                                                standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                    Moeller BL,
                                                                                                                                                                                                                                                                                                                                                                                            LUMINIS PTY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2001-374846/39
                                                                                                                                                                                                                                                  Sorghum bicolor.
                                                                                                                                                                                                                                                                            40200140491-A2
                                                                                                                                                                                                                                                                                                                                                                 01-DEC-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    biosynthesis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              16-JUL-2001
                                                                                                                                      06-AUG-2001
                                                                                                                                                                                                                                                                                                        07-JUN-2001.
                                                                                                           AAF90246;
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                                                                                AAF90246
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AAF87687/C
ID AAF876
XX
AC AAF8
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DT 16
XX
XX
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                                                                                                                                                                                                                                                                                                                                                                                                         (UYRO-)
                                                                                                                                                                                                                                                                                                                                                                                                                                     Hoej P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Matches
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The specification describes a ribozyme capable of inhibiting oestrogen-dependent tumour cell proliferation and having a high substrate specificity for an mRNA sequence encoding a DNA-binding domain of human oestrogen receptor. The ribozyme is free of endonuclease activity for an mRNA having a DNA binding domain of a glucocorticoid. The oestrogen receptor site-specific ribozymes are useful for cancer treatment and therapies, especially for inhibiting oestrogen-dependent tumour cell proliferation, particularly breast cancer. The present sequenc represents the critical sequence of a ribozyme of the invention, which targets the the DNA binding domain of a human oestrogen receptor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The invention relates to a construct which allows animals to be bred in captivity but renders them infertile in the wild by allowing reversible control over fertility and reproduction. The construct comprises a native
                       for inhibiting estrogen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     fertility; reproduction; gametogenesis; microinjection; infection;
oid gene; PCR primer; embryogenesis; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 promoter, a blocking DNA sequence contoured for and designed to abrogate a crucial gene's function or to cause its mis-expression, and a genetic switch to regulate controlled expression/repression of the blocker/gene knockout. The construct is useful for preventing embryogenesis or gametogenesis in animals by stably transforming an animal cell with the
an mRNA encoding a DNA-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 native
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mus musculus goosecoid exon 2 DNA amplifying exon 2 forward PCR primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ΡΜ.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Novel construct for preventing embryogenesis in animals comprises promoter, blocking DNA which abrogates function of crucial gene an genetic switch to regulate expression/repression of blocker/gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Grewe
Ribozyme having a high substrate specificity for an mRNA encodin binding domain of human estrogen receptor, useful for inhibiting dependent tumor cell proliferation, particularly breast cancer.
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                                                                                                                                                                                                                                                                                                                                                                                                        Length 21;
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                                                                                                                                                                                                                                                                                                                                                                                                                                             Indels
                                                                                                                                                                                                                                                                                                                                                            Sequence 21 BP; 7 A; 3 C; 8 G; 0 T; 3 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                 Score 14.2; DB 1;
Pred. No. 9.1e+02;
0; Mismatches 3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Whyard S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (CSIR ) COMMONWEALTH SCI & IND RES ORG.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example 13; Page 104; 241pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1571 ACTCAGGCAGGCCAGCTTT 1589
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                                                                                    Claim 4; Page 6; 49pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                   Query Match 0.8%;
Best Local Similarity 84.2%;
Matches 16; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Thresher R, Hinds L,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2001-425672/45.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     12-SEP-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAD09996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            knockout.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Patil J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          RESULT 1188
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mouse;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAD09996/
  g
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The present sequence is a primer used to sequence a polynucleotide encoding a human RecQ5 type DNA helicase. The three RecQ5 type helicases alpha, beta and gamma are formed by alternative splicing. The invention discloses the RecQ5 type DNA helicases beta and gamma, and the genes encoding them. The RecQ5 beta DNA helicase has a novel characteristic of being localised in the nucleus. It is useful as a diagnostic marker or in the treatment of diseases associated with chromosomal instability
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Ribozyme; oestrogen-dependent tumour; cell proliferation; glucocorticoid; DNA-binding domain; oestrogen receptor; cancer treatment; breast cancer;
                                                                                                                                                                                                                                                                                                                                                                                                      Polynucleotide encoding for RecQ5beta helicase useful for diagnosis and treatment of chromosomal instability.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Critical sequence of a ribozyme targeting the oestrogen receptor.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 14.2; DB 1; Length 21; Pred. No. 9.1e+02;
Human; RecQ5 alpha; RecQ5 beta; RecQ5 gamma; DNA helicase; alternative splicing; chromosomal instability; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           B,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 21 BP; 6 A; 2 C; 9 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                     Nishikawa K;
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                                                                                                                                                                                                                                                                                                                     Shimamoto A, Kitao S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example 2; Page 32; 97pp; Japanese
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          823 AAGTCCCTCACCCTTGTCT 841
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                                                                                                                                                                                                                                                                            (AGEN-) AGENE RES INST CO LTD
                                                                                                                                                                                           25-AUG-2000; 2000WO-JP005757.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Deery Match
Best Local Similarity 84.4%,
Conservative
Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Roy AK, Lavrovsky Y,
                                                                                                                                                                                                                                                                                                                                                            WPI; 2001-273577/28.
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                                                                                                        WO200125425-A1.
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                                                                   sapiens.
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                                                                                                                                                                                                                                    05-OCT-1999;
                                                                                                                                                                                                                                                                                                                     Furuichi Y,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        02-APR-2001
                                                                                                                                                    12-APR-2001
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                construct stably integrates into the genome by homologous recombination, and implanting the cell into a host organism, where a whole animal develops from the implanted cell. The present sequence is a PCR primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human; single nucleotide polymorphism; SNP; sickle cell anaemia; agammaglobulinaemia; diabetes insipidus; Lesch-Nyhan syndrome; muscular dystrophy; Miskott-Aldrich syndrome; Fabry's disease; familial hypercholesterclaemia; polycystic kidney disease; cancer; hereditary spherocytosis; von Willebrand's disease; tuberous sclerosis; hereditary haemorrhagic telangiectasia; familial colonic polyposis; Ehlers-Danlos syndrome; osteogenesis imperfecta autoimmune disease; acute intermittent porphyria; inflammation; nervous system disorder; infection; rheumatoid arthritis; multiple sclerosis; diabetes; systemic lupus erythematosus; Graves disease; longevity; obesity;
                                                                                                                                                                                                     Gaps
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    construct by microinjection, transfection or infection, where the
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Pred. No. 9.1e+02;
0; Mismatches 3; Indels
                                                                                                                                                                                                   3; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            oaldness; fertility; forensic; paternity testing; ss
                                                                                                                   Sequence 21 BP; 2 A; 7 C; 7 G; 5 T; 0 U; 0 Other;
                                                                            used for amplifying mouse goosecoid exon 2 DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human single nucleotide polymorphism #398.
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                                                                                                                                                                                                                                                                          21 AGTACAGAACCGGGCCCT 3
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                                                                                                                                                         0.8%;
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                                                                                                                                                                         Local Similarity 84.2
les 16, Conservative
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(IREL/) IRELAND J S.
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component is or may be genetic, such as autoimmune diseases, inflammation, cancer, diseases of the nervous system, and infection by pathogenic microorganisms, autoimmune diseases including rheumatoid arthritis, multiple sclerosis, diabetes (insulin-dependent and non-independent), systemic lupus erythematosus and Graves disease, cancers including cancers of the bladder, bain, breast, colon, oesophagus, kidney, leukaemia, liver, lung, oral cavity, ovary, pancreas, prostate, skin, stomach and uterus, longevity, appearance (e.g., baldness, obesity), strength, speed, endurance, fertility, and susceptibility or receptivity to particular drugs or therapeutic treatments), in forensics and in paternity testing. ABK65381-ABK65841 represent human single
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     agammaglobulinaemia; diaberes insipidus; Lesch-Nyhan syndrome; muscular dystrophy; Wiskott-Aldrich syndrome; Fabry's disease; muscular dystrophy; Wiskott-Aldrich syndrome; Fabry's disease; taberohelesterolaemia; polycystic kidney disease; cancer; hereditary spherocytosis; von Willebrand's disease; tuberous sclerosis; hereditary haemorrhagic telangiectasia; familial colonic polyposis; hereditary haemorrhagic telangiectasia; familial colonic polyposis; shlers-Danlos syndrome; osteogenesis imperfecta; autoimmune disease; acute intermittent porphyria; inflammation; nervous system disease; infection; rheumatoid arthritis; multiple sclerosis; diabetes; systemic lupus erythematosus; Graves disease; longevity; obesity; baldness; fertility; forensic; paternity testing; ss.
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Pred. No. 9.1e+02;
0; Mismatches 3; Indels
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IRELAND J S.
LANDER E S.
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leukaemia; neuronal disorder; stromal disorder; blastocoelic disorder;
inflammatory disorder; immune disorder; anglogenic disorder; cytostatic;
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99US-0133472P.
99US-0131459P.
99WO-US012252.
99US-0140653P.
99US-0145698P.
99US-0145698P.
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99US-014352P.
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2000WO-US000219
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99WO-US028301
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Ouery Match
Best Local Similarity 76.29
Matches 16, Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              :666
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AUG-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          08-MAR-1999;
11-MAR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        11-MAY-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       22-JUN-1999;
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01-SEP-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              15-JUL-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             02-JUN-1999
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(GETH) GENENTECH INC

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                                                                                                                                                                                                                            The present invention relates to the isolation of novel human PRO polypeptides (AAU86128-AAU86162) and the polynucleotide sequences encoding them. The PRO polypeptides, agonists, antagonists or anti-PRO antibodies are useful for treating benign or malignant tumours (e.g. renal, kidney, bladder, breast, etc), leuksemias and lymphoid malignancies, other disorders such as neuronal, glial, astrocytal, hypothalamic, glandular, macrophagal, stromal and blastocoelic disorders, inflammatory, immune and angiogenic disorders. The polynucleotide sequences are also useful in gene therapy. The present sequence represents a PCR primer used in the methods of the present invention
                                                                                                                     Thirty five nucleic acids encoding PRO polypeptides, useful for treating benign or malignant tumors, leukemias and lymphoid malignancies, inflammatory, angiogenic and immunologic disorders.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
           Hillan KJ;
Stone DM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0.8%; Score 14.2; DB 1; Length 21;
84.2%; Pred. No. 9.1e+02;
tive 0; Mismatches 3; Indels
             Gurney AL,
Smith V,
                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 21 BP; 6 A; 5 C; 8 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human polymorphism associated DNA sequence #47.
 ooddard A, Godowski PJ,
Pan J, Pitti RM, Roy MA,
Wood WI;
                                                                                                                                                                                                   Example 24; Page 136; 302pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       507 GGGCTACCTGGAGAGCTG 525
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2 GGACGACCAGGAGAAGCTG 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ABS60153 standard; DNA; 21 BP.
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23-JAN-2001; 2001US-0263678P.
02-MAR-2001; 2001US-0273037P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Best Local Similarity 84.2
Matches 16, Conservative
                                                                                        WPI; 2002-205567/26.
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               Ashkenazi AJ,
                                    Marsters SA,
Watanabe CK,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ABS60153;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RESULT 1192
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Perrone MH; Ma-Edmonds M, Zerba KE, Tsuchihashi Z, Hui L, Swanson BN, Powell JR;

WPI; 2002-619265/66.

New isolated nucleic acid with at least one polymorphic position, useful for detecting, diagnosing and treating disorders such as angioedema, cancer, viral, bacterial or fungal infection, cardiovascular and autoimmune diseases.

Disclosure; Page 706; 977pp; English.

The invention relates to an isolated nucleic acid from a human gene encoding aminopeptidase P (XPNEP2), bradykinin receptor B1 (RDKB1), cleaterase inhibitor (CIMH), kallikrein cachytinin receptor B2 (BDKB2), angiotensin converting enzyme 1 (KLKI), bradykinin receptor B2 (BDKB2), angiotensin converting enzyme 2 (ACE2) or protease inhibitor 4 (P14), comprising at least one composition Also included are (1) a probe that hybridises to a polymorphic position as provided in the detailed summary of single nuclectide polymorphisms comprising additional 5 and 3' flanking genomic compaining the sample from one or more individuals and determining the nucleic acid sequence, (2) analysing (M1) at least one nucleic acid sample comprising obtaining the sample from one or more polymorphic positions in a gene cochaining the sample from one or more polymorphic positions in a gene computed of a protein selected from the group above; (3) constructing (M2) chaplotypes using the genes comprising grouping at least two nucleic acids; (4) identifying (M3) an individual at risk of developing a disorder cusing the polymorphic data; (5) a library of nucleic acids, each of which comprises one or more polymorphic positions within a gene encoding a human protein selected from the group above; and (6) genotyping (M4) and composition with a known data set. The genes, (M1, M2, M3 and M4) condicated present in at least one polymorphic position, and compositions are useful for detecting, diagnosing treating, various disorders such as angioedeame and diseases which involve angiogenesis like haemangiomas, tumours, sarcomas, Crohn's cherenting various disorders such as angioedeame and diseases with hypertension, heart failure, myocacular diseases like angina pectoris, chypertrophy, vascular diseases, aneurysm, embolism, thrombosis, coronary contrivitive various gangia and/or atherisming anterior and composition with an enries and composition, heart failure, myocacular diseases inflammance. polynucleotides are also useful for chromosome identification. Antibodies against the proteins may be utilised for immunophenotyping of cell lines hypersensitivity reactions, sepsis, autoimmune diseases, inflammatory arthritis, cancer, wounds, viral, bacterial or fungal infection, Chronic obstructive pulmonary disease (COPD) and enterocolitis (many other diseases and disorders are listed in the specification). The and biological samples. The present sequence is included in the sequence listing but is not referred to anywhere else in the specification

Sequence 21 BP; 7 A; 5 C; 5 G; 4 T; 0 U; 0 Other;

Gaps .; 0 Score 14.2; DB 1; Length 21; Pred. No. 9.1e+02; 0; Mismatches 3; Indels Query Match 0.8%; Best Local Similarity 84.2%; Matches 16; Conservative

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1246 TICCGTATCTTAGGAACCC 1264 rrcagrererrregaacce 3

qq

ABS60250 standard; DNA; 21 RESULT 1193 ABS60250
ID ABS.
XX
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DT 05-1
XX
XX
KW Amil

(first entry) 05-NOV-2002

ABS60250;

Human polymorphism associated DNA sequence #144.

Aminopeptidase P; XPNEP2; bradykinin receptor B1; ds; BDKRB1; tachykinin receptor B1; TACR1; C1 esterase inhibitor; C1NH; kallikrein 1;

angiotensin converting enzyme 2; ACE2; protease inhibitor 4; PI4; angiotensin converting enzyme 2; ACE2; protease inhibitor 4; PI4; polymorphism; haemangioma; tumour; sarcoma; Crohn's disease; trachoma; cardiovascular disease; angina pectoris; hypertension; heart failure; myocardial infarction; ventricular hypertrophy; vascular disease; aneurysm; embolism; thrombosis; coronary artery disease; angioedaema; arteriosclerosis; atherosclerosis; hypersensitivity; sepsis; autoimmune disease; inflammatory arthritis; cancer; wound; viral infection; bacterial infection; fungal infection; COPD; chronic obstructive pulmonary disease; enterocolitis. bradykinin receptor B2; BDKRB2; gene therapy;

WO200261131-A2.

38-AUG-2002.

03-DEC-2001; 2001WO-US047235.

04-DEC-2000; 2000US-0251015P. 23-JAN-2001; 2001US-0263678P. 02-MAR-2001; 2001US-0273037P.

(BRIM) BRISTOL-MYERS SQUIBB CO. (TSUC/) TSUCHIHASHI Z. (TSUC/) TSUCHIF (HUIL/) HUI L.

Perrone MH;

Zerba KE, Ma-Edmonds M,

, Hui L, Powell JR Swanson BN,

rsuchihashi Z,

New isolated nucleic acid with at least one polymorphic position, us for detecting, diagnosing and treating disorders such as angioedema, cancer, viral, bacterial or fungal infection, cardiovascular and WPI; 2002-619265/66.

Disclosure; Page 722; 977pp; English

autoimmune diseases.

The invention relates to an isolated nucleic acid from a human gene encoding aminopeptidase P (KPNBP2), bradykinin receptor B1 (RDKRB1), cachykinin receptor B2 (RAKR), cleaterase inhibitor (CINMI), kallikrein (KIKII), bradykinin receptor B2 (BDKRB2), angiotensin converting enzyme 1 (ACE2) or protease inhibitor 4 (PI4), comprising at least one composition. Also included are (1) a probe that hybridises to a polymorphic position as provided in the detailed summary of single nucleotide polymorphisms comprising additional 5' and 3' flanking genomic comprising (MI) at least one nucleic acid sample comprising obtaining the sample from one or more individuals and determining the nucleic acid sequence at one or more polymorphic positions in a gene comprising grouping at least two nucleic acids encoding a protein selected from the group above; (3) constructing (M2) haplotypes using the genes comprising grouping at least two nucleic acids encoding a protein of an ACE inhibitor and/or vasopetidase inhibitor using the polymorphic data; (5) a library of nucleic acids, eacto of which comprises one or more polymorphic positions within a gene encoding a human protein selected from the group above; and (6) genotyping (M4) an individual comprising obtaining a nucleic acid sample, determining the culdividual comprising obtaining a nucleic acid sample, determining the culdivalual composition with a known data set. The genes, (M1, M2, M3 and M4) and composition with a known data set. The genes, (M1, M2, M3 and M4) confocial proverenting various disorders such as angiodeemes and diseases which involve angiogenesis like haemangiomas, tumours, sarcomas, crohn's disease, trachomas, and cardiovascular diseases and composition, heart failure, myocardial infarction, ventricular cornary contary anterposis, and disease, anterprosis, and contains and hypersensitivity reactions, sepsis, autoimmune diseases, inflammatory arthritis, cancer, wounds, viral, bacterial or fungal infection, Chronic obstructive pulmonary disease (COPD) and enterocolitis (many other diseases and disorders are listed in the specification). The polynucleotides are also useful for chromosome identification. Antibodies against the proteins may be utilised for immunophenotyping of cell lines

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                                                                                                                                                                                                                                                                                                         Aminopeptidase P; XPNEP2; bradykinin receptor B1; ds; BDKRB1; tachykinin receptor B1; TACR1; Cl esterase inhibitor; CINH; kallikrein 1; KLK1; bradykinin receptor B2; BDKRB2; gene therapy; angiotensin converting enzyme 2; ACB2; protease inhibitor 4; P14;
and biological samples. The present sequence is included in the sequence listing but is not referred to anywhere else in the specification
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New isolated nucleic acid with at least one polymorphic position, useful
                                                                                                                                                                                                                                                                                                                                                          polymorphism; haemangioma; tumour; sarcoma; Crohn's disease; trachoma; cardiovascular disease; angina pectoris; hypertension; heart failure; myocardial infarction; ventricular hypertrophy; vascular disease; aneurysm; embolism; thrombosis; coronary artery disease; angioedaema; arteriosclerosis; atherosclerosis; hypersensitivity; sepsis; autoimmune disease; inflammatory arthritis; cancer; wound; viral infection; bacterial infection; fungal infection; COPD;
                                                                                        Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           for detecting, diagnosing and treating disorders such as angloedema, cancer, viral, bacterial or fungal infection, cardiovascular and autoimmune diseases.
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                                                              Score 14.2; DB 1; Length 21; Pred. No. 9.1e+02;
                                                                                       Indels
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                                                                                        3;
                                      Sequence 21 BP; 5 A; 4 C; 7 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Ma-Edmonds M,
                                                                                                                                                                                                                                                                                 Human polymorphism associated DNA sequence #143.
                                                                                        0; Mismatches
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                                                                                                                  1537 AAGGAGGCCAGCCTTCGGT 1555
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(TSUC/) TSUCHIHASHI Z.
(HUIL/) HUI L.
                                                                                                                                           20
                                                                                                                                                                                                         ABS60249 standard; DNA; 21 BP.
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23-JAN-2001; 2001US-0263678P.
02-MAR-2001; 2001US-0273037P.
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                                                               0.8%;
                                                                                                                                          2 AAGGIGGACAGICTICGGI
                                                                                                                                                                                                                                                         (first entry)
                                                                                        16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Z, Hui L,
Powell JR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2002-619265/66
                                                               Query Match
Best Local Similarity
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23-JAN-2001;
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                                                                                                                                                                                                                                   ABS60249;
                                                                                                                                                                                RESULT 1194
                                                                                          Matches
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conclaiming the sample from one or more individuals and determining the obtaining the sample from one or more polymorphic positions in a genee encoding a protein selected from the group above: (3) constructing (M2) haplotypes using the genes comprising grouping at least two nucleic acids concluded at the group above: (3) constructing (M2) conditions the polymorphic data; (5) a library of developing a disorder upon administration of an ACE inhibitor and/or vasopeptidase inhibitor using the polymorphic data; (5) a library of nucleic acids, each of which comprises one or more polymorphic positions within a gene encoding a nucleotide present in at least one polymorphic position, and (6) genotyping (M4) an individual comprising obtaining a nucleic acid sample, determining the nucleotide present in at least one polymorphic position, and comparing at least one positions are useful for detecting, diagnosting, treating, and compositions are useful for detecting, diagnosting, treating, and compositions are useful for detecting, diagnosting, treating, preventing various disorders such as angloedaema and diseases which involventing various disorders such as angloedaema and diseases which they are argined and cardiovascular diseases like angina pectoris, hypertension, heart failure, myocardial infarction, ventricular context, areachomas, and cardiovascular diseases like angina pectoris, hypertension, heart failure, myocardial infarction, thrombosis, coronary artery disease, arteriosclerosis and/or atherosclerosis, and hypersensitivity reactions, sepsis, autoimmune diseases, inflammatory contextive pulmonary disease (COPD) and entercoolitis (many other diseases and disorders are listed in the specification). The polymorleotides are also useful for chromosome identification, Antibodies and also useful for chromosome identification and biological samples. The present immunophenotyping of cell lines and biological samples. The present immunophenotyping of the sequence of listing but is not referred to anywhere else in the speci
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 14.2; DB 1;
Pred. No. 9.1e+02;
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23-JAN-2001; 2001US-0263678P.
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Best Local Similarity
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02-MAR-2001; 2001US-0273037P

Perrone MH

Ma-Edmonds M,

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cuccining the sample in the or more polymorphic obstitions in a gene encoding a protein selected from the group above; (3) constructing (M2) and individual at least two nucleic acids; (4) identifying (M3) an individual at least two nucleic acids; (4) identifying (M3) an individual at least two nucleic acids; (8) allorary of nucleic acids, each of which comprises one or more polymorphic positions within a gene encoding a human protein selected from the group above; and (6) genotyping (M4) an individual comprising obtaining a nucleic acid sample, determining the nucleotide present in at least one polymorphic position, and comparing at least one position with a known data set. The genes, (M1, M2, M3 and M4) and compositions are useful for detecting, diagnosing, treating, and compositions are useful for detecting, diagnosing, treating, and compositions are useful for detecting, the agine pectoris, involve angiogenesis like haemangiomas, tumours, sarcomas, Crohn's converting various disorders such as angioedame and diseases which involve angiogenesis like haemangiomas, tumours, sarcomas, Crohn's disease, trachomas, and cardiovascular diseases like angina pectoris, hypertrophy, vascular diseases, aneurysm, embolism, thrombosis, coronary artery disease, arteriosclerosis aneurysm, embolism, thrombosis, coronary artery disease, arteriosclerosis and/or atherosclerosis, and arthritis, cancer, wounds, viral, bacterial or fungal infection, Chronic obstructive pulmonary disease (CoPD) and entercoolitis (many other diseases and disorders are listed in the specification). The polymuclocides are also useful for chromosome identification. Antibodies against the proteins may be utilised for immunophenotyping of cell lines and biological samples. The present sequence is included in the sequence is included in the sequence of listing but is not referred to anywhere else in the specification
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       encoding aminopeptidase P (XPNEP2), bradykinin receptor B1 (BDKRB1), tachykinin receptor B1 (TACR1), C1 esterase inhibitor (CINH), kallikrein 1 (KLK1), bradykinin receptor B2 (BDKRB2), angiotensin converting enzyme 2 (ACE2) or procease inhibitor 4 (P14), comprising at least one polymorphic position. Also included are (1) a probe that hybridises to a polymorphic position as provided in the detailed summary of single nucleotide polymorphisms comprising additional 5' and 3' flanking genomic sequence; (2) analysing (M1) at least one nucleic acid sample comprising obtaining the sample from one or more individuals and determining the
                                                                                                                                                                                                                                                                                                    New isolated nucleic acid with at least one polymorphic position, us for detecting, diagnosing and treating disorders such as angioedema, cancer, viral, bacterial or fungal infection, cardiovascular and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The invention relates to an isolated nucleic acid from a human gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 21 BP; 7 A; 5 C; 5 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                            Disclosure; Page 876; 977pp; English.
                                                                                                                                                                 Zerba KE,
                                                 (BRIM ) BRISTOL-MYERS SQUIBB CO. (TSUC/) TSUCHIHASHI Z.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    rrcagrercrrregaaccc 3
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                                                                                                                                                               Tsuchihashi Z, Hui L,
Swanson BN, Powell JR,
                                                                                                                                                                                                                                                                                                                                                             cancer, viral, bacte
autoimmune diseases.
                                                                                                                                                                                                                                                WPI; 2002-619265/66
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Matches 16; Conserv
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0.8%; Score 14.2; DB 1; Length 21; 84.2%; Pred. No. 9.1e+02; tive 0; Mismatches 3; Indels 1246 TICCGIAICTIAGGAACCC 1264

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Gaps

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The invention comprises a mutant form of the human aquaporin 5 (AQP5) gene. Aquaporin (AQP) is a family of water channel proteins, through which water is transported into and out of cells - ten types of mammalian AQP have been identified so far. The invention also comprises an oligonucleotide (OGN) chip having 902 oligonucleotide primer sequences and a cDNA chip comprising one or more sequences from the human AQP5 gene. The mutant AQP5 gene is useful for diagnosing cancer (i.e lung cancer). The OGN chip is useful for detecting mutations and polymorphisms in AQP5 and the cDNA chip is useful for analysis of gene expression. The present DNA sequence represents a human aquaporin (AQP) gene PCR primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Novel aquaporin 5 gene mutant useful for diagnosing lung, stomach, colon, prostate, or head or neck cancer.
                                              Human; 88; PCR; primer; aquaporin; AQP5; AQP; water channel protein; oligonucleotide chip; OGN chip; cDNA chip; lung cancer; mutation detection; polymorphism detection; gene expression.
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34.2%; Pred. No. 9.1e+02;
Lve 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                       Um T,
                                                                                                                                                                                                                                                                                                                                                                     Moon C, Moon Y, Kim B, Kim D, Shin C, Kim H, Song S;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 21 BP; 3 A; 6 C; 7 G; 5 T; 0 U; 0 Other;
              Human aquaporin 5 (AQP5) gene PCR primer 3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example 2; Page 148; 154pp; English.
                                                                                                                                                                                                                             10-SEP-2001; 2001WO-KR001528.
                                                                                                                                                                                                                                                              09-SEP-2000; 2000KR-00053821.
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                                                                                                                                                                                                                                                                                             (GOOD-) GOODGENE INC.
(MOON/) MOON W.
(MOON/) MOON C.
                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2002-393847/42.
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                                                                                                                                                        WO200220787-A1.
                                                                                                                       Homo sapiens
                                                                                                                                                                                          14-MAR-2002.
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                                                                                                                                                                                                                                                                                                                                                                     Moon W,
Song M,
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Human, 88; PCR; primer, aquaporin, AQP5; AQP; water channel protein; oligonuclectide chip; OGN chip; cDNA chip; lung cancer; mutation detection; polymorphism detection; gene expression. Human aquaporin 5 (AQP5) gene PCR primer 1. 1036 TITGGCCTGGCCCGAGCCA 1054 21 ВЪ TTTGGCCTGGCCATAGGCA ABQ61241 standard; DNA; 21 (first entry) WO200220787-A1. 03-OCT-2002 Homo sapiens ABQ61241; RESULT 1197 ABQ61241 à

14-MAR-2002.

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Gaps

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The invention comprises a mutant form of the human aquaporin 5 (AQP5) gene. Aquaporin (AQP) is a family of water channel proteins, through which water is transported into and out of cells - ten types of mammalian AQP have been identified so far. The invention also comprises an oligonuclectide (OGN) chip having 902 oligonuclectide primer sequences and a cDNA chip comprising one or more sequences from the human AQP5 gene. The mutant AQP5 gene is useful for diagnosing cancer (i.e lung cancer). The OGN chip is useful for detecting mutations and polymorphisms in AQP5 and the cDNA chip is useful for analysis of gene expression. The present DNA sequence represents a human aquaporin (AQP) gene PCR primer
Novel aquaporin 5 gene mutant useful for diagnosing lung, stomach, colon, prostate, or head or neck cancer.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human; chromosome 1p36-35; chromosome 21q22.1; genetic analysis; genome;
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                                                                                                                                                                                                                                                                                                                                                                                                   0.8%; Score 14.2; DB 1; Length 21; 84.2%; Pred. No. 9.1e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                  3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human chromosome 1p36-35 PCR primer SEQ ID NO:301
                                                                                                                                                                                                                                                                                                                                                          Sequence 21 BP; 3 A; 6 C; 7 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                  0; Mismatches
                                                                     Disclosure; Page 148; 154pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1036 TTTGGCCTGGCCCGAGCCA 1054
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(GENO-) GENOTEX YG.
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Matches 16; Conservative
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ABL43257
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                                                                                                                                                                                                                                                                                              Novel aquaporin 5 gene mutant useful for diagnosing lung, stomach, colon, prostate, or head or neck cancer.
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                                                                                                                                                                                    Kim H;
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                                                                                                                                                                                    Shin C,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 21 BP; 3 A; 6 C; 7 G; 5 T; 0 U; 0 Other
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                                                                                                                                                                                    Kim D,
                                                                                                                                                                                                                                                                                                                                                                 Example 1; Page 146; 154pp; English.
                                                                                                                                                                                    Kim B,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Kim B,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1036 TTTGGCCTGGCCCGAGCCA 1054
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ABQ61247 standard; DNA; 21 BP
      10-SEP-2001; 2001WO-KR001528
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                                                  09-SEP-2000; 2000KR-00053821
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                                                                                                                                                                                      Moon Y,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (first entry)
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Best Local Similarity 84.2
Matches 16; Conservative
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                                                                                                                                                                                                        Kim H, Song S;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (GOOD-) GOODGENE INC.
                                                                                              GOODGENE INC.
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                                                                                                                                                                                                                                                       WPI; 2002-393847/42.
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                                                                                                                                                                                 Moon C,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Kim H,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (MOON/) MOON W.
(MOON/) MOON C.
                                                                                                                 MOON W.
                                                                                                                                          MOON/) MOON C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WO200220787-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Homo sapiens,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     03-OCT-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          14-MAR-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ABQ61247;
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Song M,
                                                                                                                                                                                    Moon W,
Song M,
                                                                                              (GOOD-)
                                                                                                                     MOON/)
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8 Dp

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The present invention describes a ribozyme comprising Hu MT-Ia Rz, Hu MT-IC Rz, Rz1-2, or Rz4-9

(see ABM88812 to ABM88819. The ribozymes have cytostatic activity. The ribozymes are targeted to metallothionein (MT) and so are metallothionein concercibles in human cancer cells, for inhibiting tumour growth, and for chancing the effectiveness of chemotherapy or radiation therapy against cancer cells. The ribozyme-based methods for treating cancer, from the present invention, offer the following advantages over conventional artisense-based methods of limiting metallothionein production in target cells: (1) ribozymes destroy metallothionein-encoding mRNAs rather than core be recycled to degrade multiple mRNA molecules; (3) a ribozyme need con have perfect complementarity with a target mRNA to destroy the RNA; can be recycled to degrade multiple mRNA molecules; (3) a ribozyme need con have perfect complementarity with a target mRNA to destroy the RNA; can did (4) a single ribozyme can be designed to deetroy several related conventional antisense molecule can be designed to be effective against conventional antisense molecule can be designed to be effective against various mRNAs. ABN88819 to ABN88870 represent sequences used in the
                                                                                                                                                                                                                                                                      .
0
are detected from the amplified products; (h) the clones in the multiwell plates are specified from the detected result; and (i) the clones are reconstituted as the positions on the chromosome and arrayed. The microarray is useful for gene analysis. ABL4957 to ABL45322 represent PCR primers for human chromosome 1p36-35 DNA, and ABL45321 to ABL45634 represent PCR primers for human chromosome 21q22.1, which are
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New ribozymes directed against metallothionein mRNAs, useful for inducing apoptosis in human cancer cells, for inhibiting tumor growth and for enhancing the effectiveness of chemotherapy or radiation therapy against
                                                                                                                                                                                                                                                                      Gaps
                                                                                                                                                                                                                                                                      ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Apoptosis-inducing ribozyme; hammerhead ribozyme; ribozyme; MT;
                                                                                                                                                                                                                           Length 21;
                                                                                                                                                                                                                                                                   3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Rat metallothionein MT-II target sequence SEQ ID NO:47.
                                                                                                                                   specifically claimed for use in the present invention
                                                                                                                                                                                Sequence 21 BP; 7 A; 5 C; 4 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                        Score 14.2; DB 1;
Pred. No. 9.1e+02;
                                                                                                                                                                                                                                                                   0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (UYMA-) UNIV MASSACHUSETTS MEDICAL CENT.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 metallothionein; cancer; tumour; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 2; Fig 2B; 63pp; English.
                                                                                                                                                                                                                                                                                                            597 CTTTGGGAAACTGGAGACC 615
                                                                                                                                                                                                                                                                                                                                                     CATTCAGAACTGGAGACC 21
                                                                                                                                                                                                                        0.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         31-OCT-2001; 2001WO-US046062.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    31-OCT-2000; 2000US-0244709P
                                                                                                                                                                                                                                                                                                                                                                                                                                                               ABN88844 standard; RNA; 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   21-AUG-2002 (first entry)
                                                                                                                                                                                                                                                                   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Lee K, Lau K, Ho S;
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                                                                                                                                                                                                                        Query Match
Best Local Similarity
Matches 16; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WO200236740-A2.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ABN88844;
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Human; ds; cytochrome P450 A1; CYP4501A1; UGT2B4; MDR1;

cytochrome P450 A2; CYP4501A2; cytochrome P450 02B; CYP45002E1; LTF;

adrenergic receptor beta1; ADBR1; aryl hydrocarbon; AHR; MRB3; NRIIZ;

adrenergic receptor nuclear translocator; ARNT; cathepsin S; CTSS;

cyclooxgenase 2; COX2; diazepam binding inhibitor; DB1; haematological;

poxide hydroxylase 2; EPRK2; 5-lipoxygenase activating protein; FLAP;

W pluathione-S-transferase 12; GST12; histamine-N-methyl transferase;

HNMT; kallikrein 2; KLK2; nicotinamide-N-methyl transferase;

W MADPH quinone oxidoreductase 21; NQO2; sulfotransferase thermolabile; STM;

W UDP-glucuronosyl transferase 284; UDP-glucuronosyl transferase 2B7;

W UDP-glucuronosyl transferase; UGT2B15; urokinase receptor;

W multidrug resistance 1; lactotransferrin; orphan nuclear receptor;

W multidrug resistance associated protein 3; cancer; prostate;

M central nervons system; pulmonary; immunological; SNP;

Central nervons system; pulmonary; immunological; SNP;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Isolated nucleic acid molecules having polymorphisms in known human genes e.g. cytochrome p450 and cathepsin S useful as genetic linkage markers for locating, identifying and characterizing the genes responsible for
                                                                                                                                Gaps
                                                                                                                                .
                                                                                    Length 21;
                                                                                  Query Match 0.8%; Score 14.2; DB 1; Length 2 Best Local Similarity 68.4%; Pred. No. 9.1e+02; Matches 13; Conservative 3; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                            Human epoxide hydroylase 2 polymorphic sequence #77.
                                           Sequence 21 BP; 6 A; 4 C; 8 G; 0 T; 3 U; 0 Other;
exemplification of the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 10; Page 119; 714pp; English.
                                                                                                                                                                     1167 GGGCTGCATCTTCTATGAG 1185
                                                                                                                                                                                           2 GGGCUGCAUCUGCAAAGAG 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              single nucleotide polymorphism
                                                                                                                                                                                                                                                                                                                    BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               28-NOV-2001; 2001WO-US044838.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        28-NOV-2000; 2000US-00724389.
                                                                                                                                                                                                                                                                                                                  ABS97586 standard; DNA; 21
                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        disorder-related traits.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (DNAS-) DNA SCI LAB INC.
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                                                                                                                                                                                                                                                                                                                                                            ABS97586;
                                                                                                                                                                                                                                                                             RESULT 1201
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currensferase (NNWT), NADPH quinone oxidoreductase 2 (NQO2),
sulfotransferase thermolabile (STM), UDP-glucuronosyl transferase 2B4
(UGT2B4), UDP-glucuronosyl transferase 2B7 (UGT2B7), UDP-glucuronosyl
transferase (UGT2B15), uroxinase receptor (UPA), multidrug resistance 1
(MDR1), lactotransferrin (LTF), multidrug resistance associated protein 3
(MRP2), orphan nuclear receptor (NRR12), or acetylcholine muscarinic
creceptor 1, 2, 3, 4, or 5 (CHWR1, CHWR2, CHWR3, CHWR4 or CHMR5) sequence.
The polymorphisms in the human genes cited in the invention are useful as genetic linkage markers for locating and characterising the genes that are responsible for specific traits within the genome and eventually identifying the genes responsible for a variety of disorder-related are responsible for specific traits within the genome and eventually craries as a result of their e.g., overexpression, constitutive and/or treating the disorders. The nucleic acid molecules comprising the captures contained in CYPP4501A1, CYPP4501A1, CYPP4501E1, CYPP4501B1, C polymorphic DNA sequence of the invention

Sequence 21 BP; 3 A; 9 C; 4 G; 5 T; 0 U; 0 Other;

0.8%; Score 14.2; DB 1; Length 21; 34.2%; Pred. No. 9.1e+02; very 0; Mismatches 3; Indels 1644 GCTGGAGGGATGCCACACC 1662 Query Match 0.8%; Best Local Similarity 84.2%; Matches 16; Conservative à

21 GGTGGAAGGATGGCACACC 3

d

Gaps ; 0

RESULT 1202

ABS97587 standard; DNA; 21 BP ABS97587,

ABS97587;

(first entry) 23-DEC-2002

Human epoxide hydroylase 2 polymorphic sequence #78.

KW Human, ds, cytochrome P450 A1, CYP4501A1, UGT2B4; MDR1,

KW Cytochrome P450 A2, CYP4501A2; cytochrome P450 02E; CYP4501E1; LTF;

KW adrenergic receptor beta1; ADBR1; aryl hydrocarbon, AHR; MRP3; NRI12;

KW Adrenergic receptor muclear translocator; ARNT; cathepsin S; CTSS;

KW ADPRICATION TECEPTOR TRANSLOCATOR; ARNT; CATHEPSIN S; CTSS;

KW Cycloxygenase 2; EDHX2; 5-lipoxygenase activating protein; FLAP;

Gutathicone-S-transferase 12; GST12; histamine-N-methyl transferase;

KW NADPH quinone oxidoreductase 2 NQO2; sulfortransferase thermolabile; STM;

KW THAT; kallikrein 2; KLKZ; nicotinamide-N-methyl transferase 187;

UDP-glucuronosyl transferase 28; UDP-glucuronosyl transferase 287;

UGT2B7; UDP-glucuronosyl transferase; UGT2B15; urokinase receptor; uPA;

KW multidrug resistance associated protein 3; cancer; prostate;

multidrug resistance associated protein 3; cancer; prostate;

multidrug resistance associated protein 3; cancer; prostate;

cetylcholine muscarinic receptor; CHMR1; CHMR2; CHMR3; CHMR5;

alterated drug metabolism; cardiovascular function; colorectal tumour;

KW single nucleotide polymorphism.

WO200257410-A2

25-JUL-2002.

28-NOV-2001; 2001WO-US044838

28-NOV-2000; 2000US-00724389.

(DNAS-) DNA SCI LAB INC.

Guida M, Hall J;

WPI; 2002-698522/75.

Isolated nucleic acid molecules having polymorphisms in known human genes e.g. cytochrome p450 and cathepsin S useful as genetic linkage markers for locating, identifying and characterizing the genes responsible for disorder-related traits.

Example 10; Page 119; 714pp; English.

This invention relates to the sequence of an isolated nucleic acid
molecule comprising at least one base variation from that of a known
cut chuman cytochrome P450 Al (CYP4501A1), cytochrome P450 A2 (CYP4501A2),
cytochrome P450 Al (CYP4501A1), adrenergic receptor betal (ADBR1),
cytochrome P450 OZE1 (CYP4500ZE1), adrenergic receptor betal (ADBR1),
cytochrome P450 OZE1 (CYP4500ZE1), adrenergic receptor bidacepam binding
cytochrome P450 OZE1 (CYP4501A2), cytochrome P450 DEAL
(ARA), parly hydrocarbon receptor nuclear translocator
(ARA), property (CYSS), cytochromede 2 (GYC12), histamine-N-methyl
transferase (HNWT), NADPH quinone oxidoreductase 2 (NOO2),
transferase (HNWT), NADPH quinone oxidoreductase 2 (NOO2)
transferase (HNWT), nantheran 2) KLK2, nicothianine—N-methyl
transferase (HNWT), nantheran 2) KLK2, nicothianine—N-methyl
transferase (HNWT), nothiane receptor (URD2B7), UDP-glucuronosyl transferase 287 (UGT2B7), UDP-glucuronosyl transferase 287 (UGT2B7), UDP-glucuronosyl
transferase (UGT2B15), urokinase receptor (URD2B7), untilityl
creeptor 1, 2, 3, 4, or 5 (CHMR1, CHMR2) CHMR3 CHMR5) sequence
contains in the human genes cited in the invention are useful
genetic linkage markers for locating and characterising the genes that
creeptor 1, 2, 3, 4, or 5 (CHMR1, CHMR2) CHMR3 be used in disponsible for specific traits within the genome and eventually
identifying the genes responsible for a variety of disorder-related
traits as a result of their e.g., overexpression, which may be used in disponsible
contains and/or tracting the disorders. The nucleic acid molecules contained in CYP4501A1, CYP4501A2,
ARNY, BPHX2, GST12, NNMT, NQO2, NR112, STM, UGT2B4, UGT2B4, UGT2B1, ARNY, MR1 and/or MDR3 are useful to metabolism: The polymorphic sequences contai susceptibility to colorectal timours, in DBI or CHMRI for altered central nervous system function, in FLAP and HNWT for altered pulmonary, immunological to haematological function, in KLK2 for altered serine procease activity in the prostate, in LTF for altered immunological or haematological function, in CHMR3, CHMR4 or CHMR5 for altered central and peripheral nervous system function. The present sequence represents a polymorphic DNA sequence of the invention

Sequence 21 BP; 3 A; 9 C; 4 G; 5 T; 0 U; 0 Other;

Gaps .. 0.8%; Score 14.2; DB 1; Length 21; 84.2%; Pred. No. 9.1e+02; ive 0; Mismatches 3; Indels Local Similarity 84.2%; les 16; Conservative Query Match Best Loca Matches

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1644 GCTGGAGGGATGCCACACC 1662 21 gergaaagargecacacc

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ABK16378 standard; DNA; 21 RESULT 1203 ABK16378 HXXX

BP.

ABK16378;

10017621-381.rng

BP

(first entry)

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Adipose protein; ss; adp; obesity; transgenic animal; obesity; adipositas; bulimia; wasting; cachexia; eating disorder; body weight loss; cancer; infectious disease; hypogonadism; Prader-Willi syndrome; Laurence-Moon-Biedl syndrome; hypothyroidism; diabetes; Cushing's syndrome; endocrine disorder; hypothyroidism; diabetes; inflammatory bowel disease; PCR primer: ulcerative colitis; anorexia nervosa; glycogen storage disease; lipid storage disease; lipoma; liposarcoma; heart disease; hypertension; infertility; acquired immunodeficiency syndrome; AIDS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Novel nucleic acid encoding adipose polypeptide which regulates, causes
                                                                                                                                        Human adipose protein, adp, PCR primer #7.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          16-JUN-2000; 2000US-0211914P.
23-JUN-2000; 2000EP-00113049.
28-JUN-2000; 2000US-0214518P.
17-APR-2001; 2001EP-00109537.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   13-JUN-2001; 2001WO-EP006713.
    ABK16377 standard; DNA; 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Broenner G, Ciossek T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (DEVE-) DEVELOGEN AG.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2002-106464/14.
                                                                                                                                                                                                                                                                                                                                                                                                                                                         WO200196371-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                 Homo sapiens.
                                                                                           14-MAR-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      30-DEC-2001
                                              ABK16377;
       The invention relates to a nucleic acid encoding a adipose (ADP)

polypeptide which regulates, causes or contributes to obesity in an animal or a human. The polymucleotides, proteins, ant-adp antibodies, modulators of adp activity, adp antisense nucleic acids, expression vectors, adp transgenic animals are useful in the diagnosis and treatment of obesity, adiposites, bulimia, wasting (acabaxia), eating disorders and/or disorders of body weight/body mass, weight loss due to cancer or infectious diseases, genetic disorders associated with hypogonadism e.g. rader wills syndrome, haurence-Moon-Biedl syndrome, hypothyroidism, diseases, inflammatory bowel disease, ulcerative colitis, and anorexia diseases, inflammatory bowel disease, ulcerative colitis, and anorexia corvosa. They are also useful for treating disorders of body weight/mass e.g. glycogen storage diseases, and lipid storage diseases and for treating lipomas, and/or liposarcomas. The compositions are also useful to treating negative content of the process of the treating heart disease, hypertension, and infertility and for treating content of the process of the treating heart disease, hypertension, and infertility and for treating content of the present of the 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     treating conditions associated with under weight e.g. enhancing or controlling fertility, controlling weight loss in acquired immunodeficiency syndrome (AIDS) or cancer patients. The present sequence is a PCR primer used to amplify an adp nucleic acid
                                                                                                         adipositas; bulimia; wasting; cachexia; eating disorder; body weight disorder; weight loss; cancer; infectious disease; hypogonadism; Prader-Willi syndrome; Laurence-Moon-Biedl syndrome; hypothyroidism; diabetes; Cushing's syndrome; endocrine disorder; gastrointestinal diseases; inflammatory bowel disease; PCR primer; ulcerative colitis; anorexia nervosa; glycogen storage disease; lipid storage disease; lipoma; liposarcoma; heart disease; hypertension; infertility; acquired immunodeficiency syndrome; AIDS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Novel nucleic acid encoding adipose polypeptide which regulates, causes or contributes to obesity, useful for treating obesity, heart disease, hypertension, infertility, and controlling weight loss in cancer
                                                                                         Adipose protein; ss; adp; obesity; transgenic animal; obesity;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0.8%; Score 14.2; DB 1; Length 21;
84.2%; Pred. No. 9.1e+02;
tive 0; Mismatches 3; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 21 BP; 4 A; 8 C; 5 G; 4 T; 0 U; 0 Other;
                                            Human adipose protein, adp, PCR primer #8.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 1; Page 171; 188pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    23-JUN-2000; 2000EP-00113049.
28-JUN-2000; 2000US-0214518P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                       13-JUN-2001; 2001WO-EP006713
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(first entry)
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Best Local Similarity 84.2'
Matches 16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Broenner G, Ciossek T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (DEVE-) DEVELOGEN AG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2002-106464/14.
                                                                                                                                                                                                                                                                                                                                                              WO200196371-A2.
                                                                                                                                                                                                                                                                                                                      Homo sapiens
14-MAR-2002
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Haeder T, Rothe M;

Dohrmann C,

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The invention relates to a nucleic acid encoding a adipose (ADP)

Combined which regulates, causes or contributes to obesity in an animal or a human. The polymouleotides, proteins, ant-adp antibodies, modulators of adp activity, adp antisense nucleic acids, expression vectors, adp transgenic animals are useful in the diagnosis and treatment of observed in the diagnosis and treatment of of obesity, adiposites, bulimia, wasting (cachexia), eating disorders and/or disorders of body weight/body mass, weight loss due to cancer or infectious diseases, genetic disorders associated with hypogonadism e.g. Prader Willi syndrome, Laurence-Moon-Biedl syndrome, hypothyroidism, diabetes, Cushing's syndrome, endocrine disorders, gastrointessinal chisabetes, Cushing's syndrome, endocrine disorders, and anorexia nervosa. They are also useful for treating disorders of body weight/mass corporating they are also useful for treating disorders of body weight/mass corporated with under weight e.g. enhancing or treating lipomas, and/or liposarcomas. The compositions are also useful corporations associated with under weight e.g. enhancing or controlling fertility, controlling weight loss in acquired immunodeficiency syndrome (AIDS) or cancer patients. The present sequence is a PCR primer used to amplify an adp nucleic acid.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ö
or contributes to obesity, useful for treating obesity, heart disease, hypertension, infertility, and controlling weight loss in cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              .
0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0.8%; Score 14.2; DB 1; Length 21; 34.2%; Pred. No. 9.1e+02; ve 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 21 BP; 4 A; 5 C; 8 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                         Claim 1; Page 171; 188pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1029 GGCTGACTTTGGCCTGGCC 1047
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        19 GGCACACTTTCGCCTGGCC 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          84.28;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     16;
                                                                                         patients
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
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Gaps

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1029 GGCTGACTTTGGCCTGGCC 1047

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GGCACATTCGCCTGGCC

RESULT 1204 ABK16377/c

21

Mouse adipose complement related protein zsig37 primer ZC18687.

13-JAN-2003 (first entry)

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UGT1A7; uridine diphosphate-5'-glucuronosyl transferase; UGP; primer; carcinoma; inflammatory bowel disease; genetic predisposition; colon; polymorphism; UGT1A7*2; UGT1A7*3; UGT1A7*4; antitumour; cytostatic; antiinflammatory; gene therapy; diagnosis; pancreas; liver; stomach;
                                                                                                                                                                                                                                                                                                                                                         Diagnosing, and predicting risk, of carcinoma and inflammatory bowel disease, comprises detecting polymorphisms in the gene for uridine diphosphate-5'-glucuronosyl transferase.
                                                                                       Human UGT1A7 codon 11 polymorphism associated primer A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 21 BP; 2 A; 7 C; 7 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                          (MEDI-) MEDIZINISCHE HOCHSCHULE HANNOVER.
                                                                                                                                                                                                                                                                                                                                                                                                        Example 1; Page 12; 26pp; German.
                      BP.
                                                                                                                                                                                                                                               03-JAN-2002; 2002WO-DE000003.
                                                                                                                                                                                                                                                                    05-JAN-2001; 2001DE-01000238
                      ABL61474 standard; DNA; 21
                                                                 (first entry)
                                                                                                                                                                                                                                                                                                                Manns M, Strassburg C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               UGT1A7 gene
                                                                                                                                                                                                                                                                                                                                       WPI; 2002-509023/54.
                                                                                                                                                                                                   WO200253770-A2.
                                                                                                                                                          oesophagus; ss
                                                                                                                                                                              Homo sapiens.
                                                                                                                                                                                                                        11-JUL-2002.
                                                                 17-SEP-2002
                                            ABL61474;
RESULT 1205
           ABL61474
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This invention describes a novel method of predicting the risk, and/or diagnosis, of carcinoma and inflammatory bowel disease (IBD) associated with a genetic predisposition. The method comprises testing a subject's DNA for the presence of a polymorphic UGTLA7 allele (UGT = uridine diphosphate-5'-glucuronosyl transferase) that contains mutations in codons II, 129, 131 and/or 208. Polymorphic UGTLA7*2, UGTLA7*3 or carbolic characterisation of antitumour therapeutics and for examining textbolic characterisation of antitumour therapeutics and for examining texticity/carcinogenicity of potential UGTLA7 substrates. The products of the invention have cytostatic and antiinflammatory activity and are appropriate for gene therapy. The method of the invention is used for diagnosis, or assessing risk, of carcinoma, especially of the colon, indentification of subjects at risk. This sequence represents a primer used in the identification of the UGTLA7 polymorphism at codon II of the
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à
                                          0
                                          Gaps
                                          °,
0.8%; Score 14.2; DB 1; Length 21; ilarity 84.2%; Pred. No. 9.18+02; Conservative 0; Mismatches 3: Indels
                                                                          938 GIGGCCIGGCCIACIGCCA 956
                                                                                                       GTGGACTGGCCTCCTTCCA 21
        Query Match
Best Local Similarity
Matches 16; Conserv
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ABX04548 standard; DNA; 21 BP.

RESULT 1206

ABX04548,

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ABX04548

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The invention relates to promoting blood flow within the vasculature of a mammal, comprises administering to the mammal an amount of a mammal, comprises administering to the mammal an amount of a pharmaceutical formulation that comprises an adipocyte complement related protein; zsig37, having residues 26-281 of a sequence appearing as Composition, zsig37, having residues 26-281 of a sequence appearing as ABG99070. Also included is a method of pacifying damaged collagenous tissues within a mammal, comprising administering to the mammal an amount complication of the damaged collagenous tissues by inhibiting complement activation or by reducing thrombosis formulation. The method is useful in promoting blood flow within the vasculature of a mammal by reducing thrombogenic and complement activity, and in pacifying damaged collagenous surfaces (e.g. in trauma, ischaemia, reperfusion, intestinal stramman sursopaam; stroke, percutaneous transluminal angioplasty, constructory accidental vascular trauma or surgical infarction, cardiopulmonary bypass ischaemia, myocardial infarction, cantabody are useful as inhibitors of hasemostaasis and immune function, in complating wound healing, and for antimicrobial applications. The buman complement used to sequence cDNA encoding mouse zsig37
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Promoting blood flow within the vasculature of a mammal, comprises administering a pharmaceutical formulation comprising zsig37 proteins.
                                                                                          Mouse; ss; primer; adipocyte complement related protein; zsig37; chromosome 17q25.2; blood flow; vulnerary; antibacterial; vasotropic; anticoagulant; immunosuppressive; damaged collagenous tissue; complement activation; thrombosis; trauma; ischaemia; reperfusion; intestinal strangulation; cardiopulmonary bypass ischaemia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ..
0
                                                                                                                                                                                                  myocardial infarction; post-trauma vasospasm; stroke;
percutaneous transluminal angloplasty; endarterectomy;
accidental vascular trauma; surgical-induced vascular trauma;
haemostašis; wound healing; ancimicrobial.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0.8%; Score 14.2; DB 1; Length 21;
44.2%; Pred. No. 9.1e+02;
ve 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 21 BP; 5 A; 4 C; 8 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Lasser GW, Bishop PD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example 9; Col 53; 39pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  822 GAAGICCCICACCCIIGIC 840
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         21 GAAGTCCCTCTCACGTGTC 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ACD26013 standard; DNA; 21 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                17-FEB-2000; 2000US-00506855.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         99US-00253604.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      84.28;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (ZYMO ) ZYMOGENETICS INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2003-038245/03.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Best Local Similarity
                                                                                                                                                                                                                                                                                                          Mus musculus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            19-FEB-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sheppard PO,
                                                                                                                                                                                                                                                                                                                                                   US6448221-B1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               22-NOV-1999;
                                                                                                                                                                                                                                                                                                                                                                                            10-SEP-2002.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ACD26013/c
ID ACD26
XX
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ACD26013;

10017621-381.rng

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The invention relates to an antisense oligonucleotide complementary to a region of the open reading frame of human folate receptor alpha comprising a 774-bp sequence. Also included are inhibiting growth of cancer cells susceptible to growth inhibition, a ribozyme containing the antisense oligonucleotide and a vector comprising the antisense oligonucleotide. The antisense oligonucleotide is useful for preparing a composition for treating cancer of the ovary, cervix, uterus and brain. The present sequence is an antisense oligonucleotide targeting the human
                                                                                            Human; ss; antisense; folate receptor alpha; cytostatic; gene therapy; ribozyme; ovarian cancer; cervical cancer; uterine cancer; brain cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mouse, tryptase-like protein, ztryp-1; cardiovascular, cardiant, antiinflammatory, antiarthritic, antiinfertility, contraceptive, protein therapy, contractile tissue dysfunction; cardiovascular disease, inflammatory actions in heart; inflammatory bowel disease, arthritis; infertility; impotence, male reproductive dysfunction; birth control; in vitro fertilisation; birth; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                               antisense oligonucleotide, useful for preparing a composition for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mouse tryptase-like polypeptide Ztryp-1 related PCR primer #2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0.8%; Score 14.2; DB 1; Length 21;
84.2%; Pred. No. 9.1e+02;
live 0; Mismatches 3; Indels
                                                               Human Folate receptor alpha antisense oligonucleotide #8.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 21 BP; 2 A; 8 C; 5 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 10; Page 10; 23pp; English.
                                                                                                                                                                                                                                                                                                                                                                                 ĸ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1076 ACTCCAATGAGGTGGTGAC 1094
                                                                                                                                                                                                                                                                                                                                                                                Chung
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   20 ACCCCAATGAGGAGGTGGC 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     BP.
                                                                                                                                                                                                                                               11-MAR-2002; 2002US-00093523.
                                                                                                                                                                                                                                                                               09-MAR-2001; 2001US-0274249P.
                              (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    folate receptor alpha cDNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ACD25911 standard; DNA; 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                Jhaveri MS, Elwood PC,
                                                                                                                                                                                                                                                                                                            (JHAV/) JHAVERI M S.
(ELWO/) ELWOOD P C.
(CHUN/) CHUNG K.
                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2003-503577/47.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local Similarity
Matches 16; Conserv
                                                                                                                                                                               US2003050267-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                               treating cancer
                                                                                                                                                Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mus musculus
                              01-SEP-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     JS6514741-B1
                                                                                                                                                                                                                13-MAR-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   29-AUG-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ACD25911;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RESULT 1208
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The invention decsribes a new polypeptide (ZTRYPI) having a sequence comprising amino acid residues 44 (Val) - 276 (Ile), 24 (Leu) - 276 (Ile), 24 (Leu) - 276 (Ile), 04 (Mal) - 314 (Leu), 07 1 (Mer) - 314 (Leu), 07 1 (Mer) - 312 (Leu), 07 3 (Mal) - 275 (Mag), 19 (Marg) - 275 (Mag), 43 (Val) - 312 (Leu), 19 (Marg) - 312 (Leu) of a 312-amino acid Homo saptens sequence (hsp) or CC a 233 fusion polypeptide sequence. The ZTRYPI polypeptide is useful cor treating a dysfunction associated with contractile tissues), and may be used for suppressing or enhancing contractility in vivo. In particular, the ZTRYPI polypeptide is useful cortracting actions in heart), inflammatory cardiovascular disease (e.g. inflammatory actions in heart), inflammatory control in view dysfunction. The polypeptide is also useful in birth control, in vitro fertilisation, or inducing birth. This sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            chromosome 17q25.2; vascular occlusion; vasodilation;
                                                                                                                                                                                                                     New tryptase-like polypeptides (ZTRYPI), useful for treating a dysfunction associated with contractile tissues (e.g. heart), for modulating contractility, or for treating e.g. cardiovascular disease,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human; zsig37; ss; chromosome 17q25.2; vascular occlusion; vascdila adipocyte complement related protein; vascular injury; vascular reconstruction; trauma; stroke; ameuryam; plaque rupture; vasculature; diabetes; atheroscierosis; blood flow; vasorelaxant; tranquiliser; vulnerary; cerebroprotective; antiatherosclerotic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ;
0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0.8%; Score 14.2; DB 1; Length 21;
84.2%; Pred. No. 9.18+02;
Live 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 21 BP; 1 A; 9 C; 4 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human zsig37 cDNA sequencing primer #26.
                                                                                                                                                                                                                                                                                                         Example 1; Col 63-64; 40pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1195 GGCCGTCCCTCTTCCGG 1213
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2 decrercición con 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      BP
                                                                                  99US-0149563P.
                                                09-AUG-2000; 2000US-00636382.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     19-JUL-2000; 2000US-00619740.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADC01969 standard; DNA; 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first entry)
                                                                                                                  (ZYMO ) ZYMOGENETICS INC
                                                                                                                                                                                                                                                                           arthritis or infertility
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
                                                                                                                                                    Presnell SR, Taft DW;
                                                                                                                                                                                     WPI; 2003-491701/46.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Local Similarity
es 16; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 sequencing; primer.
                                                                                  18-AUG-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      18-DEC-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     JS6544946-B1
              04-FEB-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     08-APR-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADC01969;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
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Gaps ·.

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sheppard PO,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mus musculus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      19-FEB-1999;
22-NOV-1999;
                                                                                                                                                                                                                                                                                                                                                                                  18-DEC-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         31-JUL-2003
                                                                                                                                                                                                                                                                                                                                                             AAD59914;
                                                                                                                                                                                                        Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (LASS/)
(BISH/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (SHEP/)
                                                                                                                                                                                                                                                                                                                RESULT 1211
                                                                                                                                                                                                                                Matches
                                                                                                                                                                                                                                                                                                                           AAD59914/
                                                                                                                                                                                                                                                                                                                                       X & X O O O O O O O O O O O O X &
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                                                                                                                                                                                                                                                                             g
                                                                                                                                                                                                                                                                                                                                                                        ö
                                                                                                                                                                                                                                                       vascular injury is due to plaque rupture, degradation of the vasculature, complications associated with diabetes and atheroscilerosis.

Administration of the formulation promotes blood flow or elicits a vasorelaxant response. This sequence represents a primer used to sequence CDNA encoding the human zsig37 polypeptide of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      cardiant; antiinflammatory; antiasthmatic; antiarthritic; antiinfertility; contraceptive; serine protease; cancer; immune disorder; Ztrypl; inflammatory disorder; reproductive disorder; infertility; contraceptive; testicular disorder; heart disorder; asthma; arthritis; mouse; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New ztrypl gene, useful in diagnosing diseases associated with the ztrypl
gene, e.g., cancer or immune disorders.
                                                                                                                Minimizing vascular occlusion or inducing vasodilation within the vasculature of a mammal, by administering an adipocyte complement related protein, zsig37 that promotes blood flow.
                                                                                                                                                                                  The invention relates to a method for minimising vascular occlusion or inducing vasculation within a mammal; involving administering of formulation comprising an adipocyte complement related protein, zsig37. The method is useful for minimising vascular occlusion and inducing vasculation in a mammal suffering from acute vascular injury which may be due to vascular reconstruction, tranma, stroke or aneurysm. The vascular injury is due to plaque rupture, degradation of the vasculature,
                                                                                                                                                                                                                                                                                                                                                                         Gaps
                                                                                                                                                                                                                                                                                                                                                                         ..
0
                                                                                                                                                                                                                                                                                                                                                 0.8%; Score 14.2; DB 1; Length 21; 34.2%; Pred. No. 9.1e+02;
                                                                                                                                                                                                                                                                                                                                                                         Indels
                                                                                                                                                                                                                                                                                                                           Seguence 21 BP; 5 A; 4 C; 8 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mouse serine protease ztryp1 primer seq id 5.
                                                                                                                                                                                                                                                                                                                                                                         0; Mismatches
                                                                                                                                                              Example 9; SEQ ID NO 41; 44pp; English.
                                                                    PD;
                                                                      Bishop
                                                                                                                                                                                                                                                                                                                                                                                                822 GAAGTCCCTCACCCTTGTC 840
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 99US-00253604.
                                                                                                                                                                                                                                                                                                                                                                                                                       GAAGTCCCTCTCACGTGTC
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          22-NOV-1999; 99US-00444794.
17-FEB-2000; 2000US-00506855.
                                                                                                                                                                                                                                                                                                                                                              84.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADC17380 standard; DNA; 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
                                                                      Lasser GW,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (ZYMO ) ZYMOGENETICS INC.
                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                               (ZYMO ) ZYMOGENETICS INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Taft DW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2003-645495/61.
                                                                                             WPI; 2003-707011/67.
                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity
Matches 16; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   US2003119035-A1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          26-JUN-2003.
   19-FEB-1999;
                                                                    Sheppard PO,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             18-DEC-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADC17380;
                                                                                                                                                                                                                                                                                                                                                                                                                       21
                                                                                                                                                                                                                                                                                                                                                                                                                                                          RESULT 1210
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADC17380
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The invention describes a new isolated polynucleotide encoding a serine protease polypeptide comprising a sequence of amino acid residues that is 90% identical to a sequence comprising; amino acid residues 44-276, 24-276, 44-314, 24-314 or 1-314 or maino acid sequence or amino acid sequence or amino acid sequence, or 233 amino acids. 19-312 or 1-312 of the 312-amino acid sequence, or 233 amino acids. The polynucleotide is useful in diagnosing diseases associated with the ztrypl gene, e.g., cancer or immune disorders. Ztrypl proteins are useful for treating inflammatory, reproductive (e.g. infertility and contraceptive), testicular and heart disorders. They are also useful for treating asthma and arthritis. This sequence represents a primer used in the isolation and analysis of mouse
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The invention relates to a method of promoting blood flow within the vasculature of a mammal. The method involves administering an adipocyte complement related protein (Acrp) to the mammal to reduce and complement activity within the vasculature. Methods and compositions of the invention are useful in promoting blood flow within the vasculature of a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Adipocyte complement related protein, collagenous surface pacification, wound healing; tumour metastasis, gene therapy; thrombogenic, mouse;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Promoting blood flow within the vasculature of a mammal, comprising administering an adipocyte complement related protein to reduce thrombogenic and complement activity within the vasculature.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ·;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0.8%; Score 14.2; DB 1; Length 21; 84.2%; Pred. No. 9.1e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 21 BP; 1 A; 9 C; 4 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ZC18687 oligo used to identify mouse zisg37 DNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0; Mismatches
Example 1; SEQ ID NO 5; 44pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Bishop PD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example 9; Page 29; 48pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1195 GGCCGTCCCCTCTTCCGG 1213
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22-NOV-1999; 99US-00444794
17-FEB-2000; 2000US-00506855
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAD59914 standard; DNA; 21
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           serine protease ztrypl
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2003-755532/71.
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BISHOP P D.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      identifying polynucleotides and polypeptides whose expression levels correlate with compound sensitivity or resistance of cells associated with a disease state; and (3) identifying polynucleotides and polypeptides that predict compound sensitivity or resistance of cells associated with a disease state, comprising subjecting the plurality of cell lines to one or more compounds, analysing the expression pattern of a microarray of polynucleotides or polypeptides that predict the sensitivity or resistance of cells associated with a disease state by using the expression pattern of the microarrayy. The polynucleotides and
mammal, in pacifying collagenous surfaces, in modulating wound healing mediating tumour metastesis. The invention is also useful in gene therapy. The present sequence is an oligo used to identify mouse adipocyte complement related protein homologue (zsig37) DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      predictor set; protein tyrosine kinase activity modulator;
protein tyrosine kinase pathway; protein tyrosine kinase; cytostatic;
gene therapy; drug sensitivity; genetic profile; cancer; human;
                                                                                                                                                                                       Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New polynucleotides and polypeptides for predicting the activity of compounds that interact with protein tyrosine kinases and/or protein tyrosine kinase pathways.
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                                                                                                                                            0.8%; Score 14.2; DB 1; Length 21;
84.2%; Pred. No. 9.1e+02;
tive 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human src biomarker reverse PCR primer SEQ ID NO:600.
                                                                                                    Sequence 21 BP; 5 A; 4 C; 8 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Shaw P;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 2; SEQ ID NO 600; 139pp; English.
                                                                                                                                                                                                                         822 GAAGTCCCTCACCCTTGTC 840
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                                                                                                                                                                                                                                                                                                                                                                        ADD14411 standard; DNA; 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                         01-JAN-2004 (first entry)
                                                                                                                                        Query Match
Best Local Similarity 84.2'
Matches 16; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2003-636735/60.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PCR primer;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                 ADD14411;
                                                                                                                                                                                                                                                                                                                                    RESULT 1212
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The invention comprises oligonucleotides for detecting and identifying subtypes of human papilloma virus (HPV) contained in a sample. The oligonucleotides of the invention are useful for simultaneously detecting and identifying subtypes of HPVs. The present DNA sequence represents an oligonucleotide that was used in the exemplification of the invention.
polypeptides have cytostatic activities, and can be used in gene therapy. The polynucleotides and polypeptides are useful in predicting the activity of compounds that interact with protein tyrosine kinases and/or protein tyrosine kinase pathways. These may be used in determining drug genetic profiles which aid in treating diseases and disorders (e.g. cancer) based on patient response at a molecular level. The present sequence is used in the exemplification of the present invention.
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                                                                                                                                                                                                                Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   probe; human papilloma virus; HPV; detection; identification; ss;
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                                                                                                                                                                                                              3; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                       HPV detection method-related oligonucleotide Gap21-3.
                                                                                                                                               Seguence 21 BP; 2 A; 8 C; 3 G; 8 T; 0 U; 0 Other;
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Р;
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                                                                                                                                                                              Query Match 0.8%; Score 14.2; DB 1;
Best Local Similarity 84.2%; Pred. No. 9.1e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Lee
                                                                                                                                                                                                              0; Mismatches
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Pan C,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Kao Y,
                                                                                                                                                                                                                                            18 ATGGACAGGAATGCAGAGG 36
                                                                                                                                                                                                                                                                            19 ATGGAGAGAACTGCAGAGG 1
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                                                                                                                                                                                                              16; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Unidentified
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Hsu H,
                                                                                                                                                                                                                                                                                                                          RESULT 1213
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                                                                                                                                                                                                                                                                                                                                                                                                             A detecting apparatus and a detecting method for identifying the subtypes of many species of human papilloma viruses at the same time and a composition for the detection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           This invention describes a novel detecting apparatus for identifying the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         flow; adipocyte complement related protein; thrombogenic activity;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     complement activity; vasculature; cardiopilmonary bypass ischaemia; resuscitation; myocardial infarction; post-trauma vasospasm; stroke; percutaneous transluminal angioplasty; endarterectomy; accidental vascular trauma; surgical-induced vascular trauma; thrombosis formation; mouse; zsig37; ss; primer; sequencing.
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0
                                                                                                                                                         detection; human papillomavirus; HPV subtype; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          represents a PCR primer used in the method of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0.8%; Score 14.2; DB 1; Length 21; 34.2%; Pred. No. 9.1e+02; ive 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 21 BP; 5 A; 9 C; 3 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example 2.1.1; SEQ ID NO 649; 166pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mouse zsig37 sequencing primer #10.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1156 ATGTGGGGTGTGGCTGCA 1174
                                                                                                                                                                                                                                                                                                                                                 (KING--) KING CAR FOOD IND CO LTD
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                BP.
                             BP.
                                                                                                                                                                                                                                                                                    28-NOV-2001; 2001JP-00362595
                                                                                                                                                                                                                                                                                                                  04-MAY-2001; 2001TW-00110785
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   llarity 84.2%;
Conservative
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                               ADF44292 standard; DNA; 21
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                                                                                            (first entry)
                                                                                                                            HPV PCR primer GAP 21-3
                                                                                                                                                                                         Human papillomavirus.
                                                                                                                                                                                                                                                                                                                                                                               WPI; 2003-600935/57.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Local Similarity
es 16; Conserv
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                                                                                                                                                                                                                       JP2002360271-A.
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                                                                                              12-FEB-2004
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                                                             ADF44292;
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RESULT 1214
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             vasculature of a mammal comprises administering an adipocyte complement related protein in a vehicle, where the adipocyte complement related protein reduces thrombogenic and complement activity within the vasculature. The methods and compositions of the present invention are useful for diagnosing and treating damaged collagenous tissues, such as cardiopulmonary bypass ischaemia and resuscitation, myocardial infarction or post-trauma vasospasm including stroke, percutaneous transluminal angioplasty, endarterectomy, accidental vascular trauma or surgical induced vascular trauma. They can also be used in reducing thrombosis formation within the vasculature of a mammal. The present sequence is
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    promoting blood flow within the vasculature of mammals using an adipocyte complement related protein, useful for diagnosing and treating cardiopulmonary bypass ischemia, myocardial infarction, stroke and/or vascular trauma.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              cytostatic; gene therapy, PRO; PRO197; PRO207; PRO226; PRO232; PRO243; PRO256; PRO269; PRO274; PRO304; PRO339; PRO1558; PRO779; PRO1185; PRO1265; PRO5775; PRO5775; PRO5773; PRO5725; PRO202; PRO206; PRO364; PRO313; PRO542; PRO713; PRO661; PRO116; PRO156; PRO366; PRO360; PRO360; PRO360; PRO360; PRO360; PRO360; PRO580; 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The invention relates to a method of promoting blood flow within the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ·:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 21 BP; 5 A; 4 C; 8 G; 4 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 14.2; DB 1;
Pred. No. 9.1e+02;
0; Mismatches 3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                used in the exemplification of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 9; SEQ ID NO 41; 43pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                              Bishop PD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         822 GAAGICCCICACCCIIGIC 840
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                                                                                                                                             19-FEB-1999; 99US-00253604.
22-NOV-1999; 99US-00444794.
17-FEB-2000; 2000US-00506855.
                                                                                  10-SEP-2002; 2002US-00241258.
                                                                                                                                                                                                                                   17-JUL-2002; 2002US-00198695
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0.8%;
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Matches 16; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                              Lasser GW,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADJ37408 standard; DNA;
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                                                                                                                                                                                                                                                                                          (SHEP/) SHEPPARD P. (LASS/) LASSER G W. (BISH/) BISHOP P D.
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                                                                                                                                                                                                                                                                                                                                                                                                              Sheppard PD,
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                           24-APR-2003
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10017621-381.rng

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prevention and treatment of cancers, e.g. lymphoma, blastoma, sarcoma or leukemia, and as predictors of the prognosis of tumor treatment.
                                                                                New isolated PRO polypeptides, useful as targets for the diagnosis,
                                                                                                         Example 24; SEQ ID NO 127; 307pp; English.
                                              Goddard A, Godowski PJ,
Pan J, Pitti RM, Roy MA,
           31-AUG-1999; 99US-0151689P.
11-FEB-2000; 2000WO-US003565.
09-AUG-2001; 2001US-00927796.
02-AUG-2002; 2002US-00211858
                                                   Marsters SA, Pan J, I
Watanabe CK, Wood WI;
                                 (GETH ) GENENTECH INC
                                                                     WPI; 2003-901564/82.
                                             Ashkenazi AJ,
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0.8%; Score 14.2; DB 1; Length 21; 34.2%; Pred. No. 9.1e+02; ve 0; Mismatches 3; Indels Sequence 21 BP; 6 A; 5 C; 8 G; 2 T; 0 U; 0 Other; 84.2%; Matches 16; Conservative Query Match Best Local Similarity

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Gaps

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ABX99015 standard; DNA; 21 BP (first entry) 20-MAY-2003 ABX99015; RESULT 1217 ABX99015,

Human AAGA SNP analysis PCR primer, #42.

Human; PCR; primer; ss; asthma; bronchial hyperresponsiveness; airway obstruction; chronic bronchial inflammation; multifactorial disease; asthma-associated gene; AAGA; allele-specific; single nucleotide polymorphism; SNP; genetic profile; gene therapy; antisense gene therapy; adult distress respiratory syndrome; chronic obstructive pulmonary; chronic bronchitis; dyspnea.

Homo sapiens.

WO2003008640-A2

30-JAN-2003.

15-JUL-2002; 2002WO-EP007847

16-JUL-2001; 2001US-0305649P

Hillan KJ;

Stone DM;

Gurney AL, Smith V,

(NOVS) NOVARTIS AG.
(NOVS) NOVARTIS ERRINDUNGEN VERW GES MBH.
(UYWA-) UNIV WAKE FOREST HEALTH SCI.
(UYGR-) RIJKSUNIV GRONINGEN.

Bleecker ER Whittaker PA, Meyers DA, Postma DS,

WPI; 2003-239359/23.

Determining whether a subject has or is at risk of developing a disease characterized by bronchial hyperresponsiveness, comprises determining the expression or bioactivity level of an asthma-associated gene.

Example 3; Page 27; 70pp; English.

construction and primers sued to anlayse AAGA single nucleotide polymorphisms (SNPs)

Sequence 21 BP; 3 A; 5 C; 8 G; 5 T; 0 U; 0 Other;

Length 21; 3; Indels Score 14.2; DB 1; Pred. No. 9.1e+02; 0; Mismatches 0.8%; Query Match
Best Local Similarity 84.2'

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Gaps

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BP. ACD02587 standard; DNA; 21 ACD02587/

RESULT 1218

ACD02587;

31-JUL-2003 (first entry)

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zsig37 orthologue sequencing primer 2C18687 Mouse

Blood flow, vasodilation, wound repair; platelet inhibition; tumour; vascular occlusion; ischaemic reperfusion injury; microvascular repair; adipocyte complement related protein; intestinal strangulation; trauma; angioplasty; coronary artery bypass graft; endarterectomy; aneurysm; anastomosis; stroke; cardiopulmonary bypass ischaemia; inflammation; mycardial infarction; percuteneous transluminal angioplasty; infection; post-trauma vasospasm; prostatic blomaterial; fibroblast recruitment; wound retraction; mouse; zsig37; primer; ss; sequencing; PCR.

Mus musculus.

US2003022838-A1.

30-JAN-2003

25-JUN-2002; 2002US-00180762.

99US-00253604. 99US-00444794. 19-FEB-1999; 22-NOV-1999;

2000US-00506855. 2000US-00619740. 17-FEB-2000; 19-JUL-2000;

(SHEP/) SHEPPARD P O. (LASS/) LASSER G W. (BISH/) BISHOP P D.

Bishop PD; Lasser GW, Sheppard PO,

WPI; 2003-456304/43.

Promoting blood flow or inducing vasodilation within vasculature of mammal, or pacifying damaged collagenous tissues or pacifying surface of prostatic biomaterial, by administering adipocyte complement related protein.

Example 9; Page 29; 46pp; English.

vaccular injury, where the injury is due to vascular reconstruction which comprises angioplasty, coronary artery bypass graft, endarterectomy, comprises angioplasty, coronary artery bypass graft, endarterectomy, an excovascular repair or anastomosis of a vascular graft, or the injury is due to trauma, stroke or aneurysm. The method is useful for pacifying damaged collagemous tissues within a mammal, where the damaged collagemous tissues within a mammal, where the damaged collagemous tissues are due to injury associated with ischaemia intestinal collagemous tissues are due to injury associated with ischaemia intestinal repertusion. The injury associated with pre- and post-establishment of strangulation, or injury associated with pre- and post-establishment of creasitation, mycardial infarction, or post-trauma vasospasm. The post- creasitation, mycardial infarction, or post-trauma vasospasm. The post- creama vasospasm comprises stroke, percutancous transluminal angioplasty, endarterectomy accidental vascular trauma or surgical-induced vascular trauma vasospasm comprises stroke, percutancous transluminal angioplasty, endarterectomy accidental vascular trauma or surgical-induced vascular trauma vasospasm comprises stroke, percutancous transluminal angioplasty.

Crequit, fibrin or fibronectin. The method is useful for mediating wound repression in wound healing comprises reduction in the creation in the method is useful for inhibiting platelet creduction in infection. The method is useful for inhibiting platelet creduction in infection. The method is useful for inhibiting platelet creatment. The method is useful for inhibiting platelet creatment. The method is useful for inhibiting platelet creatment. The method is useful for inducing vasculating comparating comparating comparating mammal. The method is useful for mediating mediating mycoardium from reperfusion injury. The method is useful for mediating vasodilation within the vasculature of a mammal, pacifying damaged collagenous tissues or surface of prostatic biomaterial, mediathing wound repair, inhibiting platelet adhesion, activation or accretion, minimising vascular occlusion, protecting ischaemic myocardium from reperfusion protecting ischaemic myocardium from reperfusion complishing tuminour metastrasis, comprising administering adipocyte complement related protein. The method is useful for promoting blood flow within the vasculature of a mammal where the mammal suffers from acute invention relates to a method of promoting blood flow or inducing

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tumour metastasis. The present sequence represents the mouse adipocyte complement related protein zsig27 DNA orthologue sequencing primer
                                                                   Gaps
                                                                                                                                                                                                                                       cytostatic; gene therapy; vaccine; cancer; chromosome mapping;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Zhong M;
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                                               Length 21;
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                                                                   Indels
                           Sequence 21 BP; 5 A; 4 C; 8 G; 4 T; 0 U; 0 Other;
                                              Score 14.2; DB 1;
Pred. No. 9.1e+02;
0; Mismatches 3;
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                                                                                                                                                                                                                     NOVX oligonucleotide probe, SEQ ID No 221.
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                                                                                       822 GAAGICCCICACCCIIGIC 840
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2002US-0363429P.
2002US-036383P.
2002US-0372141P.
2002US-0373961P.
2002US-0373061P.
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2001US-0327893P.
2001US-0331768P.
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2002US-0358939P.
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2002US-0360830P.
2002US-0361178P.
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                                                                                                          21 GAAGICCCICICACGIGIC
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                                                                                                                                                                                                                                                                                                                                                                              2001US-0305011P.
2001US-0305262P.
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2001US-0306085P.
2001US-0307536P.
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2001US-0312203P.
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2001US-0323948P.
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                                                 0.8%;
                                                                                                                                                           ADM47388 standard; DNA; 21
                                                                                                                                                                                                 (first entry)
                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (CURA-) CURAGEN CORP
                                                           Sest Local Similarity
Matches 16; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Shenoy SG,
                                                                                                                                                                                                                                                                                        WO2003083039-A2.
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30-JUL-2001; 2
14-AUG-2001; 2
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05-MAR-2002;
12-MAR-2002;
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24-JUL-2001;
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21-SEP-2001;
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                                                                                                                                                                                                  03-JUN-2004
                                                                                                                                                                                                                                                                      Unidentified
                                                                                                                                                                                                                                                                                                                                                                                        13-JUL-2001;
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                                                                                                                                                                                ADM47388;
                                                  Query Match
                                                                                                                                         RESULT 1219
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Hillan KJ; Stone DM;

Gurney AL, Smith V,

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neoplastic tumour; lung; colon; breast; prostate; rectal; cervical; liver; gene therapy; Human; as; primer; PCR.
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                                                                     Example C; SEQ ID NO 221; 433pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                           Human PRO polypeptide PCR primer #35.
                                                                                                                                                                                                                                                                                                                                   350 TGGGGTCTGATGGGGAGAG 368
                                                                                                                                                                                                                                                                                                                                            |||||| || || || |||||||||| TGGGGGCTTATAGGGAGAG 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         02-AUG-2002; 2002US-00210951.
                                                                                                                                                                                                                                                                                                                                                                                   ADG68332 standard; DNA; 21
                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
                                                                                                                                                                                                                                                                                                                      Conservative
                                 WPI; 2003-812538/76.
                                                                                                                                                                                                                                                                                                              Local Similarity
nes 16; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            US2003170228-A1
      Shimkets RA,
Catterton E,
                                                                                                                                                                                                                                                             pathologicaî
                                                                                                                                                                                                                                                                                                                                                                                                                                                               Homo sapiens
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             Padigaru M,
                     Pena CEA,
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The invention relates to an isolated antibody that binds to a PRO polypeptide. The methods and compositions of the present invention are useful for diagnosting, prognosticating and/or treating neoplastic tumours, such as lung, colon, breast, prostate, rectal, cervical and liver tumours. The PRO polypeptides are also useful as molecular weight markers, or for chromosome identification. The PRO genes are useful as hybridisation probes, or for screening libraries of Human cDNA, genomic DNA or mRNA. The PRO genes may also be used in gene therapy, particularly for replacing a defective gene. The present sequence is used in the exemplification of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New isolated antibodies binding PRO polypeptides, useful for diagnosing, prognosticating and/or treating neoplastic tumors, such as lung, colon, breast, prostate, rectal, cervical and liver tumors.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 21 BP; 6 A; 5 C; 8 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                               Joudard A, Godowski PJ, Pan J, Pitti RM, Roy MA, Wood WI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example 24; SEQ ID NO 127; 308pp; English
                                 31-AUG-1999; 99US-0151689P.
11-FEB-2000; 2000MO-US003565.
09-AUG-2001; 2001US-00927796.
                                                                                                                                                                           (GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2004-020650/02.
                                                                                                                                                                                                                                                      Ashkenazi AJ,
                                                                                                                                                                                                                                                                                    Marsters SA,
Watanabe CK,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                         The invention relates to a novel isolated polypeptide, designated NoVX.

The novel polypeptide comprises a sequence comprising 109-1671 amino acids to the mature form; a sequence that is at least 954 identical to compensative substitutions in the 109-1671 amino acid polypeptide; or a sequence comprising one or more composition; a method for determining the presence or amount of the composition; a method for determining the presence or amount of the polypeptide or nucleic acid molecule in a sample, determining the composition; a method for determining the presence or amount of the polypeptide or nucleic acid molecule in a sample, determining the composition; a method for of expression of the polypeptide in a first mammalian subject; identification to, a disease associated with the altered levels of nucleic acid or of expression of the polypeptide in a first mammalian subject; identification of an agent that binds to the polypeptide; dentification of a potential therapeutic agent for treating a pathology related to aberrant expression or physiological interactions of the polypeptide; a method for modulating the activity of the polypeptide; a method for modulating the activity of the polypeptide; treating a pathological state in a mammal; an isolated nucleic acid molecule; a coll comprising the nucleic acid have cytostatic acid and a method for producing the polypeptide. The NoVX polypeptide can be used in create a vaccine. The polypeptide is composition for treating or preventing a composition for treating or preventing a composition for treating or preventing a composition for preventing or preventing a composition for treating or preventing a composition for treating or preventing a composition of care a vaccine. The polypeptide is nathology as mannal and a method for predate a vaccine. The polypeptide is not the polypeptide in a mammal and a method for predate a vaccine. The polypeptide is not the polypeptide and a method for predate a vaccine. The polypeptide is a composition for treating or predated in a man
                                                                                                                                                                                                                                                                                           or
                                 Gerlach VL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     pathological state in a mammal, e.g., cancer, or for chromosome mapping. This polynucleotide sequence represents a probe used in the exemplification of the invention.
                                                                                                    Boldog FL;
                                                                                                                                                                                                                                                                            New NOVX polypeptide, useful for preparing a composition for treating preventing e.g. cancer or for chromosome mapping.
Edinger SR, Hjart
                      E, Spytek KA, Miller CE, Edinger SR, Hjalt T, A, Taupjer RJ, Anderson DW, Guo K, Baumgartne. Peyman JA, Smithson G, Casman SJ, Voss EZ, Chapoval A, Rastelli L, Kekuda R, Vernet CAM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 21 BP; 4 A; 10 C; 2 G; 5 T; 0 U; 0 Other;
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                         Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                          Novel polynucleotide useful for PCR amplification along with two DNA fragment from another set of sequences, or for detecting single
                         0
0.8%; Score 14.2; DB 1; Length 21; 84.2%; Pred. No. 9.18+02; ive 0; Mismatches 3; Indels
                                                                                                                                                                                                                             human; single nucleotide polymorphism; SNP; ss; primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 2; SEQ ID NO 7301; 2627pp; Japanese
                                                                                                                                                                                                                                                                                                                                                                          (KAGA-) KAGAKU GLJUTSU SHINKO JIGYODAN
                                                                                                                                                                                                                                                                                                                                                                                                                                                  nucleotide polymorphism in human gene.
                                               507 GGGCTACCTGGAGAGCTG 525
                                                                      2 GGACGACCAGGAGAAGCTG 20
                                                                                                                                                                                                      Primer of the invention #3992.
                                                                                                                                   BP
                                                                                                                                                                                                                                                                                                                            08-MAR-2002; 2002JP-00064373.
                                                                                                                                                                                                                                                                                                                                                  08-MAR-2002; 2002JP-00064373.
                                                                                                                                  ADK98272 standard; DNA; 21
                                                                                                                                                                                (first entry)
                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2004-093977/10.
           Local Similarity
Les 16; Conserv
                                                                                                                                                                                                                                                                            JP2003259875-A.
                                                                                                                                                                                06-MAY-2004
                                                                                                                                                                                                                                                                                                    16-SEP-2003.
                                                                                                                                                                                                                                                     Synthetic.
                                                                                                                                                        ADK98272;
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0;

Gaps

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The present invention relates to a polynucleotide isolated from a human

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Primer ZC18687 used to generate mouse zsig37 DNA.
                                                                                                                                                                                     Sheppard PD,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     15-JUL-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       22-APR-2004.
                                                                                       22-JAN-2004.
                              primer; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADM94143;
                                                                                                                                                                                                                                                                                                                                                                                                                                             822
                                                                                                                                                                                                                                                                                                                                                                                                                                                                21
                                                                                                                                                                                                                                                          a mammal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             RESULT 1224
                                                  Mus sp.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADM94143,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                g
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ·.
                                                                                        .
0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The present invention relates to a polynucleotide isolated from a human gene and is useful for detecting a single nucleotide polymorphism in a fumman gene or for diagnosing of disease. The invention enables the detection of a single nucleotide polymorphism in a human gene. The present sequence represents a primer of the invention.
                                                                                        Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                  Novel polynucleotide useful for PCR amplification along with two DNA fragment from another set of sequences, or for detecting single
 gene and is useful for detecting a single nucleotide polymorphism inman gene or for diagnosing of disease. The invention enables the detection of a single nucleotide polymorphism in a human gene. The present sequence represents a primer of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           .
0
                                                                                        0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 14.2; DB 1; Length 21; Pred. No. 9.1e+02; 0; Mismatches 3; Indels
                                                                   Score 14.2; DB 1; Length 21; Pred. No. 9.1e+02; 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                         single nucleotide polymorphism; SNP; ss; primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 21 BP; 6 A; 8 C; 3 G; 4 T; 0 U; 0 Other;
                                                  G; 6 T; 0 U; 0 Other,
                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 2; SEQ ID NO 5964; 2627pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                             (KAGA-) KAGAKU GIJUTSU SHINKO JIGYODAN
                                                                                                                                                                                                                                                                                                                                                                                                                                       nucleotide polymorphism in human gene.
                                                                                                             1692 CCCTGCTTACTCTCTGCCT 1710
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GACTIGAAGAIGGGGICIG 358
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ..
                                                                                                                              20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       BP.
                                                                                                                                                                              BP
                                                                                                                                                                                                                                      Primer of the invention #2655.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match

0.8%;
Best Local Similarity 84.2%;
Matches 16; Conservative 0
                                                                     ch 0.8%;
11 Similarity 84.2%;
16; Conservative C
                                                                                                                              cacreceraercrerecer
                                                                                                                                                                                                                                                                                                                                                           08-MAR-2002; 2002JP-00064373
                                                                                                                                                                                                                                                                                                                                       08-MAR-2002; 2002JP-00064373
                                                    Sequence 21 BP; 2 A; 7 C; 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADJ96243 standard; DNA; 21
                                                                                                                                                                              ADK96935 standard; DNA; 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
                                                                                                                                                                                                                     (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2004-093977/10.
                                                                       Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                JP2003259875-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     06-MAY-2004
                                                                                                                                                                                                                     06-MAY-2004
                                                                                                                                                                                                                                                                                                                    16-SEP-2003
                                                                                                                                                                                                                                                                               Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              340
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADJ96243
                                                                                                                                                                                                 ADK96935;
                                                                                                                                                                                                                                                            human;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              RESULT 1223
                                                                                                                                                             RESULT 1222
                                                                                          Matches
                                                                                                                                                                     ADK96935/
ID ADK9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADJ96243/
                                                                                                                                                                                        XXXXXX
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    55555x8
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Use of adipocyte complement related protein for promoting blood flow within the vasculature, pacifying damaged collagenous tissues, pacifying the surface of a prostatic biomaterial or mediating a wound repair within
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The present invention relates to polynucleotides and polypeptide molecules for use asinhibitors in haemostasis and immune function. The invention is useful promoting blood flow, pacifying damaged collagenous tissues, pacifying the surface of a prostatic biomaterial and mediating a wound repair within a mammal. The invention is also useful in preventing infection at the wound site. The present sequence is a primer used to to generate mouse zsig37 DNA. The primer is used in the exemplification of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  nucleic acid amplification; primer; PCR; detection; chromosome aberration; lymphoproliferative disorder; ss.
Haemostasis; immune function; wound repair; infection; zsig37; mouse;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               °;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 21 BP; 5 A; 4 C; 8 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 14.2; DB 1;
Pred. No. 9.1e+02;
0; Mismatches 3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example 9; SEQ ID NO 41; 42pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Bishop PD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TCRD gene related Vdelta4 primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              840
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                gaagrecerereacerere
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GAAGTCCCTCACCCTTGTC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0.8%;
                                                                                                                                                                                                                                                               17-JUL-2002; 2002US-00198695.
                                                                                                                                                                                                                                                                                                                     17-JUL-2002; 2002US-00198695.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity 84.2
Matches 16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Lasser GW,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADM94143 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                       (SHEP/) SHEPPARD P D. (LASS/) LASSER G W. (BISH/) BISHOP P D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2004-132060/13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WO2004033728-A2
                                                                                                                                                 US2004014650-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  the invention.
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Morgan GJ;

San Miquel JF;

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The present invention describes a set of nucleic amplification primers capable of amplifying a VH-JH or DH-JH IGH, VK-JK or VK/intron-Kde IGK, Valambda-llambda IGL, Vbeta-Jbeta TCRB or Dbeta-Deta TCRB, VY-JY TCRG, Vdelta-Jdelta, Ddelta-Ddelta or Vdelta-Ddelta TCRB rearrangement comprises a forward primer and a reverse primer. Also described: (1) a nucleic acid amplification assay, preferably a PCR or multiplex PCR or wiltiplex PCR or viriatron-Kde IGK, Vlambda-Jlambda IGL, Vbeta-Jbeta TCRB or Dbeta-CTC or VK/intron-Kde IGK, Vlambda-Jlambda IGL, Vbeta-Jbeta TCRB or Dbeta-CTC or VK/intron-Kde IGK, Vlambda-Jlambda IGL, Vbeta-Ddelta or Vdelta-Ddelta CTCRD rearrangement; (3) detecting chromosomal translocation (11;14) (BCLG-CTCB) carrangement; (3) detecting chromosomal translocation (11;14) (BCLG-CTCB) carrangement; (5) assessing clonal rearrangements and/or chromosome aberrations; and (6) a xit for the detecting at least one rearrangement comprising the set of primers. The new set of nucleic amplification primers capable of amplifying a VH-JH or DH-JH IGH, VK-JK or VK/intron-CTCRC, Vdelta-Jdelta-Ddelta Ddelta-Ddelta TCRB, vary TCRG, Vdelta-Jdelta-Ddelta or Vdelta-Ddelta TCRB, rearrangement are useful in diagnosing lymphoproliferative disorders. The present correcting a very invention.
                                                                                                                                                                                                                                                       New set of nucleic amplification primers comprising a forward primer a reverse primer and capable of amplifying a rearrangement, useful in diagnosing lymphoproliferative disorders.
                                                                                                Van Dongen JJM, Langerak AW, Schuuring EMD, San Miguel Garzia Sanz R, Parreira A, Smith JL, Lavender FL, Morg
Evans PAS, Kneba M, Hummel M, Macintyre EA, Bastard C;
                                                                                                                                                                                                                                                                                                                                                             Claim 9; Fig 9B; 121pp; English.
                     (UYRO-) UNIV ROTTERDAM ERASMUS (DAVI/) DAVI F B L.
                                                                                                                                                                                                      WPI; 2004-364878/34.
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Sequence 21 BP; 10 A; 5 C; 4 G; 2 T; 0 U; 0 Other;

Query Match 0.8%; Score 14.2; DB 1; Length 21; Best Local Similarity 84.2%; Pred. No. 9.1e+02; Matches 16; Conservative 0; Mismatches 3; Indels 1399 CIGITGCAGTITGAGGGTC 1417

21 crerrecarrirecreere 3

AAT55032 standard; RNA; 15 (revised) 25-MAR-2003 18-APR-1997 AAT55032; RESULT 1225

Human relA hammerhead ribozyme target sequence (nt. position 630) (first entry)

Enzymatic nucleic acid; ribozyme; trans cleavage; inhibition; intercellular adhesion molecule; rel A; tumour necrosis factor; intercellular adhesion molecule; rel A; tumour necrosis factor; INF-alpha; respiratory syncytial virus; RSV, bcr-abl; oncogene; translocation; chronic myelogenous leukaemia; CML; cancer; Philadelphia chromosome; inflammation; autoimmune disease; atherosclerosis; myocardial infarction; stroke, restenosis; transplant rejection; rheumatoid arthritis; psoriasis; myocardial; Kawasaki disease; septic shock; HIV; human immunodeficiency virus; acquired immune deficiency syndrome; AIDS;

Homo sapiens

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940S-00224483.
940S-00227958.
940S-00228041.
940S-00245736.
940S-00291932.
                                                                   9405-00292520
9405-00293520
9405-003030303
9405-00311486
9405-00311486
9405-00316771
9405-00316771
9405-00334847
9405-00334847
9405-00334847
9405-00334847
                                                                                                                                      95US-00380734
WO9523225-A2
                23-FEB-1995;
        31-AUG-1995
                                                   18-MAY-1994
                                                            15-AUG-1994
                                                                     17-AUG-1994
                                                                                       23-SEP-1994;
                                                                                               28-SEP-1994;
                                                                                                   03-OCT-1994)
                                                                                                                04-NOV-1994;
                                                                                                                     10-NOV-1994;
                                      07-APR-1994
                                               15-APR-1994
                                                                              02-SEP-1994
                                                                                 08-SEP-1994
                                                                                          23-SEP-1994
                                          15-APR-1994
                                                                          19-AUG-1994
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(RIBO-) RIBOZYME PHARM INC.

Stinchcomb DT, Chowrira B, Direnzo A, Draper KG, Dudycz LW; Grimm S, Karpeisky A, Kisich K, Matulic-Adamic J, Mcswiggen JA; Modak A, Pavco P, Belgleman L, Sullivan SM, Sweedler D, Thompson JD; Tracz D, Usman N, Wincott FE, Woolf T;

WPI; 1995-351090/45.

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Gaps

.,

Ribozymes having modified bases and methods for producing them - for use in inhibiting disease related genes.

Claim 2; Page 228; 407pp; English.

The present sequence represents a preferred target sequence for an enzymatic nucleic acid (i.e. a ribozyme) which cleaves relA mRNA at the nuclectide base position indicated in the DE line. The relA gene product is a subunit of the transcriptional regulator NF-kappaB and is implicated specifically in the induction of inflammatory responses. Regions of the mRNA that do not form secondary folding structures and that contain protential hammerhead and hairpin ribozyme cleavage sites were identified by computer analysis. Ribozymes directed against these mRNA sequences were designed and synthesised with modifications that improve their nuclease reastance. The ribozymes are designed to cleave the target sequences and thereby inhibit relA expression, making them potentially useful for treating rheumatoid arthritis, restenosis and asthma as well as for increasing tolerance to transplanted tissues. The potential immunosuppressive properties of a ribozyme cleaves relA mRNA means that uses are limited to local delivery, acute indications or ex vivo treatment. (Updated on 25-MAR-2003 to correct PI field.)

Sequence 15 BP; 4 A; 5 C; 1 G; 0 T; 5 U; 0 Other; Query Match

0.8%; Score 14; DB 1; Length 15; 71.4%; Pred. No. 7.4e+02; 0; Indels 4; Mismatches 71.48; 538 CCCATCTTTGACAA 551 Local Similarity 71.4 nes 10; Conservative Matches Best

.; 0

Gaps

.; 0

1 CCCAUCUUUGACAA 14

8 g BP.

(first entry)

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Antisense therapy; antiproliferative; antinflammatory; antipsoriatic; cytostatic; dermatological; cardiant; virucide; ophthalmological; keloid; skin discorder; Insulin-like Growth Factor I receptor; IGF-1; pityriasis; IGF binding protein; IGFBP-2; IGFBP3; inflammation; psoriasis; pilatis; growth factor mediated cell proliferation; ichthyosis; serborrhoea; ruba; keratosis; neophasia; scleroderma; wart; skin cancer; sclerotic disease; hyperneovascular condition; hyperplasia; kidney disease; neobascular condition; hyperplasia; kidney disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Ameliorating the effects of a disorder, e.g. psoriasis, by administering UV (ultra-violet) treatment (optional) and an antisense nucleic acid that inhibits or reduces growth factor mediated cell proliferation and/or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          21-JUN-2000; 2000WO-AU000693.
                                                                                                                                            IGF-I oligonucleotide #1576
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Wraight CJ, Werther GA,
AAF50616 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2001-041421/05.
                                                                                                                                                                                                                                                                                                                                                                                                                                                             WO200078341-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           inflammation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        21-JUN-1999;
                                                                                             30-MAR-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          28-DEC-2000.
                                               AAF50616;
skin disorders. The method comprises contacting the skin with an antisense oligonucleotide, (for Insulin-like Growth Factor [IGF]-1 receptor, IGF binding protein [IGFB]-2 or IGFBB3), which is capable of inhibiting or reducing growth factor mediated cell proliferation, inflammation and/or other disorders. The present sequence is an oligonucleotides of the present invention (see AAFA5151 and AAF45153-F45161). The method is useful for ameliorating the effects of psoriasis, ichthyosis, pityriasis, ruba, pilaris, serborrhoea, keloids, keratosis, neoplasias, scleroderma, warts, benign growths, cancers of the skin, a hoperneovascular condition such as neovascular condition of the retina, brain or skin, growth factor-mediated malignancies, other sclerotic disease, kidney disease, hyperproliferation of the inside of blood vessels or any other hyperplasia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Ameliorating the effects of a disorder, e.g. psoriasis, by administering UV (ultra-violet) treatment (optional) and an antisense nucleic acid that inhibits or reduces growth factor mediated cell proliferation and/or
                                                                                                                                                                                                                                                                                    Antisense therapy; antiproliferative; antiinflammatory; antipsoriatic; cytostatic; dermatological; cardiant; virucide; ophthalmological; keloid; skin discorder; Insulin-like Growth Factor I receptor; IGF-1; pityriasis; IGF binding protein; IGFBP-2; IGFBP3; inflammation; psoriasis; pilaris; growth factor mediated cell proliferation; ichthyosis; serborrhoea; ruba; keratosis; neophasia; scleroderma; wart; skin cancer; sclerotic disease; hyperneovascular condition; hyperplasia; kidney disease; neobascular condition; hyperplasia; kidney disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          present invention relates to a method for ameliorating the effects of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0.8%; Score 14; DB 1; Length 15; 0.00.0%; Pred. No. 7.4e+02; ve 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 15 BP; 2 A; 8 C; 3 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Edmondson SR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (MURD-) MURDOCH CHILDRENS RES INST.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example 8; Page 71; 201pp; English.
                                                                                                    BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            99US-0140345P
                                                                                                                                                                                                                                               IGF-I oligonucleotide #1580
                                                                                                    AAF50620 standard; DNA; 15
                                                                                                                                                                                               (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity 100.
Matches 14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Werther GA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2001-041421/05
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WO200078341-A1,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                inflammation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               21-JUN-1999;
                                                                                                                                                                                                 30-MAR-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Wraight CJ,
                                                                                                                                                    AAF50620;
                                                    RESULT 1226
                                                                              AAF5062(
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Edmondson SR;

(MURD-) MURDOCH CHILDRENS RES INST.

99US-0140345P.

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                                              The present invention relates to a method for ameliorating the effects of skin disorders. The method comprises contacting the skin with an antisense oligonuclectide, (for Insulin-like Growth Factor [IGF]-1 receptor, IGF binding protein [IGFB]-2 or IGFBP3), which is capable of inhibiting or reducing growth factor mediated cell proliferation, inflammation and/or other disorders. The present sequence is an oligonuclectide which can be used to design the antisense.
                                                                                                                                                                                                                                         oligomuclectides of the present invention (see AAF45151 and AAF45153-F45161). The method is useful for ameliorating the effects of psoriasis, ichthyosis, pityriasis, ruba, pilaris, serborkhea, keloids, keratosis, neoplasias, scleroderma, warts, benign growths, cancers of the skin, a hyperneovascular condition such as a neovascular condition of the retina, brain or skin, growth factor-mediated malignancies, other sclerotic disease, kidney disease, hyperproliferation of the inside of blood
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 14; DB 1; Length 15;
Pred. No. 7.4e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 15 BP; 1 A; 8 C; 4 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  .,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mismatches
Example 8; Page 71; 201pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0.5°,
100.0%; Pre
                                                                                                                                                                                                                                                                                                                                                                                                                                                    vessels or any other hyperplasia
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1100 GGTACCGGCCCCCT 1113
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ABX04015 standard; DNA; 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     14;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Local
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Matches
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Gaps 0;

100.08; Pre

1103 ACCGGCCCCTGAC 1116

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ACCGGCCCCTGAC 14

RESULT 1227 AAF50616

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Claim 10; Page 29; 58pp; German.
                                                                           13-MAR-2001; 2001DE-02010013.
                                                                                    13-MAR-2001; 2001DE-01012348.
13-MAR-2001; 2001DE-02010013.
                                                                                                                                                                                                                                                      183 CATAGACAAGACCA 196
                                                                                                                                                                                                                                                                                         ADM76115 standard; DNA; 15
09-JAN-2003 (first entry)
                                                                                                                                                                                                                                                                                                            03-JUN-2004 (first entry)
                                                                                                                                                                                                                                                               CATAGACAAGACCA 1
                                                                                                                                                                                                                                            Conservative
                                  diabetic symptom; ss
                                                                                                              WPI; 2001-657777/76.
                                                                                                                                                                                                                                Query Match
Best Local Similarity
                                                                                                    (ROET/) ROETGER A.
                                                                                                                                                                                                              the invention
                                                      DE20110013-U1.
                                                                                                                                                                                                                                                                                                                                                                     Homo sapiens
                                                                 18-OCT-2001
                                                                                                                                                                                                                                           14;
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                                                                                                                                                                                                                                                                                                   ADM76115;
                                                                                                                                                                                                                                                                              RESULT 1229
                                                                                                                                                                                                                                           Matches
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50.7% homology to the human EphA7 expressed and its gene is located on chromosome 1. Ephrin receptors are overexpressed in various cancers and it has been found that inhibition of NEPHA expression promotes apoptosis. The invention also relates to the NEPHA transcriptional control (promoter) region (ADM76037); recombinant vectors and host cells control (promoter) region (ADM76037); recombinant vectors and host cells comprising the NEPHA promoter compounds which inhibit or activate transcription of the NEPHA gene; and pharmaceutical compositions comprising an activator. The antisense oligonucleotide or a transcriptional inhibitor or activator. The antisense oligonucleotides and modulators of NEPHA transcription are useful for inducing apoptosis for the treatment and/or prevention of cancers in which NEPHA is overexpressed such as lung cancer, ovarian cancer, breast cancer, cavical cancer. Second transcription in terpresent transcription factor binding sites within the transcriptional
                                                                                                                                                                                                                                                                                                               Novel antisense oligonucleotide useful as anticancer agent for preventing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Vascular endothelial growth factor receptor; VEGF receptor; flt-1; flk-1; KDR; hammerhead ribozyme; hairpin ribozyme; cleavage; tumour angiogenesis; psoriasis; rheumatoid arthritis; ocular disease; fms-like tyrosine kinase 1; kinase insert domain containing receptor; foetal liver kinase 1; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                     The invention relates to antisense oligonuclectides (ADM76030 and ADM76031) targeted to the human NEPHA gene (ADM76029), which encodes a novel brain-derived ephrin receptor (ADM76028). The NEPHA protein has
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mouse flt-1 VEGF receptor hammerhead ribozyme substrate #456.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 14; DB 1; Length 15;
Pred. No. 7.4e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0; Indels
                                                                                                                                                                                                                                                                                                                                          cancer e.g. lung cancer, stomach cancer, breast cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 15 BP; 4 A; 3 C; 6 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mismatches
                                                                                                                                                                                                                                                                                                                                                                                        Example 2; Page 21; 38pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 control region of the NEPHA gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity 100.0%; Pr
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            BP
                                                                                                                                                      05-APR-2002; 2002JP-00103497.
                                                                                                      05-APR-2002; 2002JP-00103497.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1645 CTGGAGGGATGCCA 1658
                                                                                                                                                                                                         (TAKE ) TAKEDA CHEM IND LID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAX74928 standard; RNA; 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
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                                                                                                                                                                                                                                                            WPI; 2004-038434/04.
JP2003289876-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          W09715662-A2
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                                                   14-OCT-2003
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                                                                                              Detection; probe, diagnosis; oral disease; paradontitis; caries; therapy; polymorphism; virulence factor; antibiotic resistance gene; prognosis; oral infection; detection; pathogen; coronary heart disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        This invention describes a novel nucleotide carrier with probes used for diagnosis of oral diseases, particularly paradontitis, but also caries, especially to identify genetic predisposition (as indicated by polymorphisms) to disease and to identify causative microorganisms or their associated virulence factors and antibiotic resistance genes, e.g. for selection of therapy and for prognosis. They are also useful for research into oral infections. The carriers allow simultaneous detection of both host and pathogen parameters, providing quickly and simply an individual's paradontitis profile, including detection of pathogens that are associated with increased risk of coronary heart diseases and/or approach on the supportunistic pathogens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Oligonucleotide array, useful for diagnosing oral diseases, particularly paradontitis, carries human or microbial reference sequences.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ABX03870-ABX04044 represent DNA fragments used to illustrate the method
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human; NEPHA; ephrin receptor; brain; chromosome 1; apoptosis; drug screening; antisense therapy; gene therapy; cancer; tumour; lung cancer; ovarian cancer; breast cancer; cervical cancer; prostate cancer; bladder cancer; schomach cancer; colorectal cancer; cytostatic; transcriptional control region; promoter; transcription factor binding site; ds.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 14; DB 1; Length 15; Pred. No. 7.4e+02;
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                                             Resistance genes mefA & mefE DNA fragment.
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100.0%; Pre-
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Gaps

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96WO-US017480.

25-OCT-1996; 26-OCT-1995;

01-MAY-1997.

95US-0005974P.

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rheumatoid arthritis, etc., in a human patient
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                                                                                                                                                                                                                                                                                                                                         AAX74911;
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                                                                                                                                                                  The present invention describes nucleic acid molecules which modulate the synthesis, expression and/or stability of a mRNA encoding 1 or more receptors of vascular endothelial growth factor (WEGF). A patient (preferably human) having a condition associated with the level of the fims-like tyrosine kinase 1 (fl.1), kinase insert domain containing receptor (KDR) and/or foetal liver kinase 1 (flk-1) (e.g. tumour angiogenesis, coular diseases, psoriasis and rheumatoid arthritis) can be treated by administering the nucleic acid molecule or the expression vector to the patient. AAX67275 to AAX75752 represent specific examples of nucleic acid molecules from the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Nucleic acid molecule modulating VEGF receptor(s) gene expression or mRNA stability - useful for treating e.g. tumour angiogenesis, psoriasis,
                                                                                                    Nucleic acid molecule modulating VEGF receptor(s) gene expression or mRNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Vascular endothelial growth factor receptor; VBGF receptor; flt-1; flk-1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              KDR; hammerhead ribozyme; hairpin ribozyme; cleavage; tumour angiogenesis; psoriasis; rheumatoid arthritis; ocular disease; fms-like tyrosine kinase 1; kinase insert domain containing receptor;
                                                                                                                                                                                                                                                                                                                                           Gaps
                                                                                                               stability - useful for treating e.g. tumour angiogenesis, psoriasis, rheumatoid arthritis, etc., in a human patient.
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                                                                                                                                                                                                                                                                                                                     0.8%; Score 14; DB 1; Length 17;
71.4%; Pred. No. 8.3e+02;
Ive 4; Mismatches 0; Indels
                                                          Escobedo J;
                                                                                                                                                                                                                                                                                               Sequence 17 BP; 4 A; 4 C; 3 G; 0 T; 6 U; 0 Other;
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                                                          Stinchcomb D,
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                                                                                                                                                 Claim 4; Page 168; 218pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                              AAX71437 standard; RNA; 17 BP.
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96US-00584040.
    96US-00584040
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Best Local Similarity 71.4%;
Matches 10; Conservative
                                                                                                                                                                                                                                                                                                                                                                  539 CCATCTTTGACAAG 552
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                                                                                                                                                                                                                                                                                                                                                                             2 CCAUCUTUGACAAG 15
                        RIBOZYME PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     fms-like tyrosine kinase :
foetal liver kinase 1; ss
                                                           Mcswiggen J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mcswiggen
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                                                                                WPI; 1997-259017/23.
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                                     CHIRON CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         25-OCT-1996;
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11-JAN-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Homo sapiens
    11-JAN-1996;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   01-MAY-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAX71437;
                                                           Pavco P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Pavco P,
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                          (RIBO-)
                                     (CHIR )
                                                                                                                                                                                                                                                                                                                                                                                                                         RESULT 1231
                                                                                                                                                                                                                                                                                                                                                                                                                                 AAX71437
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synchesis, expression and/or stability of a mRNA encoding 1 or more receptors of vascular endothelial growth factor (WGF). A patient preferably human having a condition associated with the level of the fms-like tyrosine kinase 1 (fit-1), kinase insert domain containing receptor (KDR) and/or foetal liver kinase 1 (fik-1) (e.g. tumour anglogenesis, ocular diseases, psoriasis and rheumatoid arthritis) can be
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Nucleic acid molecule modulating VEGF receptor(s) gene expression or mRNA stability - useful for treating e.g. tumour angiogenesis, psoriasis, rheumatoid arthritis, etc., in a human patient.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The present invention describes nucleic acid molecules which modulate the synthesis, expression and/or stability of a mown
                                                                                                                                                                                                                                                                                  arthritis) can be
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Vascular endothelial growth factor receptor; VEGF receptor; flt-1; flk-1;
                                                                    present invention describes nucleic acid molecules which modulate the
                                                                                           synthesis, expression and/or stability of a mRNA encoding 1 or more receptors of vascular endothelial growth factor (VBGP). A patient (Preferably human) having a condition associated with the level of the fms-like tyrosine kinase 1 (IL1-1), kinase insert domain containing receptor (KDR) and/or foetal liver kinase 1 (flk-1) (e.g. tumour angiogenesis, coular diseases, psoriasis and rheumatoid arthritis) can be treated by administering the nucleic acid molecule or the expression vector to the patient. AAX67275 to AAX7572 represent specific examples of nucleic acid molecules from the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             KDR; hammerhead ribozyme; hairpin ribozyme; cleavage;
tumour angiogenesis; psoriasis; rheumatoid arthritis; ocular disease;
fms-like tyrosine kinase 1; kinase insert domain containing receptor;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mouse flt-1 VEGF receptor hammerhead ribozyme substrate #439
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 14; DB 1; Length 17; Pred. No. 8.3e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Seguence 17 BP; 4 A; 4 C; 6 G; 0 T; 3 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 4; Page 168; 218pp; English.
Claim 4; Page 110; 218pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 GGAGAAGUCCCUCA 14
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Best Local Similarity
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AAV97498 standard; RNA; 17

RESULT 1234

AAV97498/

(first entry)

17-MAR-1999

AAV97498;

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Vascular endothelial growth factor receptor; VEGF receptor; flt-1; flk-1; KDR; hammerhead ribozyme; hairpin ribozyme; cleavage; tumour anglogenesis; psoriasis; rheumatoid arthritis; ocular disease; fms-like tyrosine kinase 1; kinase insert domain containing receptor;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Nucleic acid molecule modulating VEGF receptor(s) gene expression or mRNA stability - useful for treating e.g. tumour angiogenesis, psoriasis, rheumatoid arthritis, etc., in a human patient.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The present invention describes nucleic acid molecules which modulate the synthesis, expression and/or stability of a mRNA encoding 1 or more receptors of vascular endothelial growth factor (VEGF). A patient (preferably human) having a condition associated with the level of the fms-like tyrosine kinase 1 (f1c-1), kinase insert domain containing receptor (KDR) and/or foetal liver kinase 1 (f1k-1) (e.g. tumour angiogenesis, ocular diseases, psoriasis and Theumatoid arthritis) can be treated by administering the nucleic acid molecule or the expression of nucleic acid molecules from the present invention
             vector to the patient. ÄAX67275 to AAX75752 represent specific examples of nucleic acid molecules from the present invention
                                                                                                                                     Gaps
treated by administering the nucleic acid molecule or the expression
                                                                                                                                     0;
                                                                                                                                                                                                                                                                                                                                                                             Mouse flt-1 VEGF receptor hammerhead ribozyme substrate #455.
                                                                                              0.8%; Score 14; DB 1; Length 17; 71.4%; Pred. No. 8.3e+02; cive 4; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Escopedo J;
                                                               Sequence 17 BP; 1 A; 5 C; 6 G; 0 T; 5 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 4; Page 168; 218pp; English.
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96US-00584040.
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                                                                                                                                  10; Conservative
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                                                                                                               Local Similarity
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The present invention describes enzymatic nucleic acid molecules (NAMs) which specifically cleave RNA derived from an epidermal growth factor receptor (EGF-R) gene. AAV9721 to AAV98043 and AAV98979 to AAV99000 represent specifically claimed target sequence from human EGF-R. AAV98044 to AAV98066 and AAV98867 to V9878 represent hammerhead ribozymes and hairpin ribozymes respectively for human EGF-R. The NAMs are useful for cleaving EGF-R RNA in the treatment of a condition associated with EGFR expression levels e.g. to inhibit cell proliferation in the prevention or treatment of cancers. The NAMs can also be used as diagnostic tools to examine genetic drift and mutations within diseased cells or to detect the presence of EGF-R RNA in a cell
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                                                                                                                                                                                                                                                                                                                                                                                                                                                        Enzymatic nucleic acids - which cleave RNA derived from an epidermal growth factor receptor, useful for inhibiting cell proliferation and for
                                                                                                         Human, epidermal growth factor receptor; EGFR; EGF-R; target sequence;
hammerhead ribozyme; hairpin ribozyme; inhibition; cell proliferation;
cancer; genetic drift; detection; mutation; ss.
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100.0%; Pred. No. 8.3e+02;
ive 0; Mismatches 0; Indels
                                                                            Human EGF-R target sequence nucleotide position 2416.
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                                                                                                                                                                                                                                                                                                                                                                                               Mcswiggen JA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 5; Page 73; 109pp; English.
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                                                                                                                                                                                                                                                                                                                                                             UNIV ASTON
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          treating cancers.
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Gaps

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0.8%; Score 14; DB 1; Length 17; 71.4%; Pred. No. 8.3e+02; tive 4; Mismatches 0; Indels

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    which specifically cleave RNA derived from an epidermal growth factor which specifically cleave RNA derived from an epidermal growth factor receptor (EGF-R) gene. AAV97221 to AAV98043 and AAV98979 to AAV99090 receptor (EGF-R) gene. AAV97221 to AAV98046 from human EGF-R. AAV98044 to AAV98066 and AAV98086 frow persone from human EGF-R. AAV98044 hairpin ribozymes respectively for human EGF-R. The NAMs are useful for cleaving EGF-R RNA in the treatment of a condition associated with EGFR expression levels e.g. to inhibit call proliferation in the prevention or treatment of cancers. The NAMs can also be used as diagnostic tools to examine genetic drift and mutations within diseased cells or to detect the presence of EGF-R RNA in a cell
                                                                                                                                                                                                                                                                                                                                                                                                                             Enzymatic nucleic acids - which cleave RNA derived from an epidermal growth factor receptor, useful for inhibiting cell proliferation and for treating cancers.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human; ss; antisense therapy; cytostatic; antiinflammatory; haemostatic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The present invention describes enzymatic nucleic acid molecules (NAMs) which specifically cleave RNA derived from a figure and acid.
                              Human; epidermal growth factor receptor; EGFR; EGF-R; target sequence; hammerhead ribozyme; hairpin ribozyme; inhibition; cell proliferation; cancer; genetic drift; detection; mutation; ss.
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   Human EGF-R target sequence nucleotide position 2412
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 17 BP; 4 A; 6 C; 2 G; 0 T; 5 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 5; Page 73; 109pp; English.
                                                                                                                                                                                                                                                                                                                                                                           Mcswiggen JA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ;
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100.0%;
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Best Local Similarity 100.
Matches 14; Conservative
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                                                                                                                                                                                                                                                                                                                                                                             Fell P,
                                                                                                                                                        WO9833893-A2
                                                                                                                       Homo sapiens
                                                                                                                                                                                                                               14-JAN-1998;
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The invention relates to a nucleic acid molecule which down regulates expression of a CD20 gene and a nucleic acid molecule which down regulates expression of a neurite growth inhibitor gene (NGGO). The nucleic acids may be enzymatic nucleic acids (e.g. a ribozyme or a nucleic acids may be enzymatic nucleic acid cleaving an RNA molecule DNAzyme) an Inozyme (an endolytic nucleic acid cleaving an RNA molecule of possessing an NCH motif), a C-cleaver (Cleaving RNA with a NYM motif) provided the argument RNA with an NGN triplet), a zinzyme (cleaving RNA with a VGY motif). The CD20-targetting nucleic acid is used to cleave RNA cof CD20 in the presence of a divalent cation that is preferably MG<sup>2</sup> +. CC FUTCO. The treatment may further comprise the use of one or more cof CD20. The treatment may further comprise the use of one or more cof CD20. The treatment may further comprise the use of one or more cof CD20. The treatment may further comprise the use of one or more streat lymphoma (ML), bulky low-grade or follicular NHL, lymphoma, leukaemia, HIV (human immunodeficiency virus) associated NHL, mantle-cell lymphoma (MCL), immunocytoma (IMC) small B-cell lymphocytic lymphoma, immunodeficiency virus) associated NHL, mantle-cell curgence of a divalent cation that is preferably MG<sup>2</sup> +. Furthermore, the presence of a divalent cation that is preferably MG<sup>2</sup> +. Furthermore, the creatment may further comprise the use of one or more cell and treat a patient having a condition associated with the level of therapies. In particular, the NOGO-targetting nucleic acid may be contacted with a cell to reduce NOGO activity of the central nervous system (CNG) injury and cerebrovascular accident cell and treat a patient having a condition associated with the level of therapies. In particular, the NOGO-targetting nucleic acid may be used to therapies. In particular, and injury and cerebrovascular accident central nervous system (CNG) injury and cerebrovascular accident central network a dissase, dementia, multiple sclerosis (MS), chemot
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ö
Parkinson's digease; ataxia; Huntington's disease;
Creutzfeldt-Jakob disease; muscular dystrophy; neurodegenerative disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Nucleic acid molecules, e.g., enzymatic nucleic acids and antisense constructs, which down regulate expression of a CD20 gene or neurite growth inhibitor gene useful for treating, e.g., lymphoma, leukemia, and central nervous system injury.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Parkinson's disease, ataxia, Huntington b anserve, edisease disease, muscular dystrophy, and/or other neurodegenerative disease
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Pred. No. 8.3e+02;
2; Mismatches 0; Indels
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28-FEB-2000; 2000US-0185516P.
06-MAR-2000; 2000US-0187128P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mcswiggen J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (CHOW/) CHOWRIRA B M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              BLATT L.
MCSWIGGEN J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2001-607195/69.
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Best Local Similarity
                                                                                                                                                                                                                                                                                  WO200159103-A2.
                                                                                                                                                      Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                  16-AUG-2001
                                                                                                                                                                                              Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Blatt L,
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ABK01785 standard; RNA; 17
                 (first entry)
                                                                                                           RIBOZYME PHARM INC
                       Human NOGO Zinzyme #107.
                                                                                                                         Blatt L, Mcswiggen J,
                                                                                                                MCSWIGGEN J.
CHOWRIRA B M.
                                                                                                                               WPI; 2001-607195/69.
                                                                            WO200159103-A2.
                                                                    sapiens.
                 12-MAR-2002
                                                                                  16-AUG-2001
                                                                      Synthetic.
            ABK01785;
                                                                                                          (RIBO-)
                                                                                                                (MCSW/)
                                                                                                             (BLAT/)
RESULT 1237
                                                                    Homo
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Human; ss; antisense therapy; cytostatic; antiinflammatory; haemostatic; cerebroprotective; nootropic; neuroprotective; antiparkinsonian; muscular; CD20; neurite growth inhibitor gene; NoGO; hammerhead ribozyme; DNAzyme; inozyme; G-cleaver; amberzyme; zinzyme; lymphoma; leukaemia; B-cell lymphoma; non-Bodgkin; s lymphoma; MIL; lymphocytic leukaemia; human immunodeficiency virus; HIV associated NHL; mantle-cell lymphoma; MCL; immunocytoma; IMC; immune thrombocytopaenia; stroke; dementia; inflammatory arthropathy, central nervous system injury; cerebrovascular accident, CVA, Alzheimer's disease; multiple sclerosis; chemotherapy-induced neuropathy; amyotrophic lateral sclerosis; ALS; parkinson's disease; ataxia, Huntington's disease; Creutzfeldt-Jakob disease; muscular dystrophy; neurodegenerative disease. Chowrira BM; 28-FEB-2000; 2000US-0185516P. 06-MAR-2000; 2000US-0187128P. 09-FEB-2001; 2001WO-US004273 11-FEB-2000; 2000US-0181797P

The invention of a CD20 gene and a mucleic acid molecule which down regulates expression of a neurite growth inhibitor gene (NOCO). The nucleic acids may be enzymatic nucleic acid molecule which down nucleic acids (e.g. a ribozyme or a DNAzyme) an Inozyme (an endolytic nucleic acid cleaving a an RNA molecule possessing an NCH motif), a G-cleaver (cleaving RNA with a NNN motif) proceed an amberzyme (cleaving RNA with an NGN triplei), a zinzyme (cleaving RNA with a VGY motif). The CD20-targetting nucleic acid is used to cleave RNA with a VGY motif). The CD20-targetting nucleic acid is used to cleave RNA cf CD20 in the presence of a divalent cation that is preferably Mg^2 2...

Furthermore, it may be contacted with a cell to reduce CD20 activity of the cell and treat a patient having a condition associated with the level cf CD20. The treatment may further comprise the use of one or more therapies. In particular, the CD20 targetting nucleic acid may be used to treat lymphoma, leukaemia, B-cell lymphoma, low-grade or follicular NHL, lymphocytic leukaemia, HIV fhuman immunodeficiency virus) associated NHL, mantle-cell CH lymphoma (MCL), immunocytoma (IMC) small B-cell lymphoma, immunocytoma (IMC) small B-cell lymphoma, thrombocytopaenia, and inflammatory arthropathy. The NOGO gene in the targetting nucleic acid is used to cleave RNA of the NOGO gene in the Nucleic acid molecules, e.g., enzymatic nucleic acids and antisense constructs, which down regulate expression of a CD20 gene or neurite growth inhibitor gene useful for treating, e.g., lymphoma, leukemia, and invention relates to a nucleic acid molecule which down regulates Claim 88; Page 97; 200pp; English. central nervous system injury.

Nucleic acid molecules, e.g., enzymatic nucleic acids and antisense constructs, which down regulate expression of a CD20 gene or neurite growth inhibitor gene useful for treating, e.g., lymphoma, leukemia, and central nervous system injury.

Chowrira BM;

Blatt L, Mcswiggen J, CHOWRIRA B M.

WPI; 2001-607195/69.

11-FEB-2000; 2000US-0181797P. 28-FEB-2000; 2000US-0185516P. 06-MAR-2000; 2000US-0187128P.

RIBOZYME PHARM INC.

RIBO-)

BLATT L. MCSWIGGEN J.

(CHOM/) (BLAT/)

09-FEB-2001; 2001WO-US004273.

WO200159103-A2.

16-AUG-2001.

sapiens.

Synthetic

presence of a divalent cation that is preferably Mg^2z^+. Furthermore, the nucleic acid may be contacted with a cell to reduce NOGO activity of the cell and treat a patient having a condition associated with the level of NOGO. The treatment may further comprise the use of one or more therapies. In particular, the NOGO-targetting nucleic acid may be used to treat central nervous system (CNS) injury and cerebrovascular accident (CVA, stroke), Alzheimer's disease, dementia, multiple sclerosis (MS), chemotherapy-induced neuropathy, anyotrophic lateral sclerosis (MS), Parkinson's disease, ataxia, Huntington's disease, Creutzfeldt Jakob disease, muscular dystrophy, and/or other neurodegenerative disease states which respond to the modulation of NOGO expression. The present ٥. Human; ss; antisense therapy; cytostatic; antiinflammatory; haemostatic; cerebroprotective; notropic; neuroprotective; antiparkinsonian; muscular; CD20; neurite growth inhibitor gene; NOGO; hammerhead ribozyme; DNAzyme; inozyme; G-cleaver; amberzyme; zinzyme; lymphoma; leukaemia; human immundeficioncy virus; HIV associated NHL; lymphocytic leukaemia; human immundeficioncy virus; HIV associated NHL; mantle-cell lymphoma; MCL; immuncytoma; IMC; immune thrombocytopaenia; stroke; dementia; inflammatory arthropathy; central nervous system injury; central accident; CVA; Alzheimer's disease; multiple sclerosis; chemotherapy-induced neuropathy; amyotrophic lateral sclerosis; ALS; parkinson's disease; ataxia, Huntington's disease; creutzfeldt-Jakob disease; muscular dystrophy; neurodegenerative disease. Gaps .; 0 Score 14; DB 1; Length 17; Pred. No. 8.3e+02; 2; Mismatches 0; Indels Sequence 17 BP; 2 A; 7 C; 6 G; 0 T; 2 U; 0 Other; sequence is a zinzyme molecule of the invention ABK00760 standard; RNA; 17 BP. 0.8%; 96 3 ccceceecucudas 16 12-MAR-2002 (first entry) Query Match
Best Local Similarity 85.79 83 cccccccccrcrcAc Human NOGO Inozyme #30. ABK00760; 1238 ABK00760 889999999999988 ò ద

Ellis JH;

Hamblin PA,

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The invention relates to a nucleic acid molecule which down regulates expression of a CD20 gene and a nucleic acid molecule which down regulates expression of a neurite growth inhibitor gene (NGGO). The regulates expression of a neurite growth inhibitor gene (NGGO). The nucleic acids may be enzymatic nucleic acids (e.g. a ribozyme or a nucleic acids may be enzymatic nucleic acid cleaving an RNA molecule DNAzyme) an Inozyme (an endolytic nucleic acid cleaving an RNA molecule possessing an NCH motif), a G-Cleaver (Cleaving RNA with a NYM motif) promise the CD20 in the presence of a divalent cation that is preferably Mg<sup>2</sup> +. Curhermore, it may be contacted with a cell to reduce CD20 activity of the real and treat a patient having a condition associated with the level of CD20. The treatment may further comprise the use of one or more therapies. In particular, the CD20 targetting nucleic acid may be used to treat lymphoma, leukaemia, HV (numan immunodeficiency virus) associated WHL, municetting nucleic acid may be used to leukaemia, HV (numan immunodeficiency virus) associated WHL, municetting nucleic acid may be contacted with a cell lymphocytic lymphocytic lymphoma, immunocytopaenia, and inflammatory arthropathy. The NOGO ene in the nucleic acid may be contacted with a cell to reduce NOGO activity of the cell and treat a patient having a condition associated with the level of the reatment may further comprise the use of one or more therapies. In particular, the NOGO-targetting nucleic acid may be used to treat central nervous system (CNS) injury and cerebrookscular acident
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (CVA, stroke), Alzheimer's disease, dementia, multiple sclerosis (MS), chemotherapy-induced neuropathy, amyotrophic lateral sclerosis (ALS), Parkinson's disease, ataxia, Huntington's disease, Creutzfeldt-Jakob disease, muscular dystrophy, and/or other neurodegenerative disease states which respond to the modulation of NOGO expression. The present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human, Grb2-related with Insert Domain, GRID, T-cell; co-stimulatory adaptor protein; tissue rejection; graft rejection;
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85.7%; Pred. No. 8.3e+02;
Live 2; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 17 BP; 2 A; 7 C; 6 G; 0 T; 2 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           sequence is an inozyme of the invention
                    88; Page 78; 200pp; English.
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Matches 12; Conservative
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                                                                                                                                                             The present invention relates to oligonucleotides that downregulate the expression of human Grb2-related with Insert Domain (GRID) gene. GRID is a T-cell co-stimulatory adaptor protein. The oligonucleotides are useful for modulating the expression of GRID, to treat conditions such as tissue/graft rejection and leukaemia. The oligonucleotides can also be administered in conjunction with other therapies such as radiation,
                                                                                                                                                                                                                                                               chemotherapy and cyclosporin treatment. The present oligonucleotide was used to illustrate the invention
                                                                  Domain
                                                                                                                                                                                                                                                                                                                                                                                        Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human, Grb2-related with Insert Domain; GRID; T-cell;
co-stimulatory adaptor protein; tissue rejection; graft rejection;
leukaemia; cytostatic; ss.
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                                                             nucleic acid(s) for regulating the Grb2-related with Insert I
(D) gene comprises using antisense and enzymatic nucleic acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human GRID hammerhead ribozyme substrate oligonucleotide #74.
                                                                                                                                                                                                                                                                                                                                                 0.8%; Score 14; DB 1; Length 17;
100.0%; Pred. No. 8.3e+02;
tive 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                    Sequence 17 BP; 5 A; 6 C; 1 G; 0 T; 5 U; 0 Other;
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Mcswiggen JA,
                                                                                                   molecules such as hammerhead ribozymes.
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                                                                                                                                     Claim 4; Page 60; 108pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ABL46441 standard; RNA; 17 BP.
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 Von Carlowitz I,
                                                                                                                                                                                                                                                                                                                                                                                                                           598 TTTGGGAAACTGGA 611
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                            14; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 leukaemia; cytostatic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2001-550088/61.
                                   WPI; 2001-550088/61.
                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity
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     Jarvis T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ABL46441;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 RESULT 1240
                                                                                       (GRID)
                                                                                                                                                                                                                                                                                                                                                                                             Matches
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PAPP-E; human; pregnancy associated plasma protein E; abortive; contraceptive; gene therapy; vaccine; pregnancy; antenatal; diagnosis; dysgenetic pregnancy; primer; ss.

06-APR-2001; 2001US-00827998.

US2002102252-A1.

01-AUG-2002.

26-MAY-2000; 2000US-0207456P.

SHANNON M E.

GUYY/) (SHAN/)

WPI; 2002-697817/75. Gu Y, Shannon ME;

Human PAPP-Ea associated 17-mer SEQ ID 541.

(first entry)

24-DEC-2002

ABS75015;

ABS75015 standard; DNA; 17

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chemotherapy and cyclosporin treatment. The present oligonucleotide was
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Domain
                                                                                                                   Gaps
                                                                                                                                                                                                                                                                                                                                                                                          Human, Grb2-related with Insert Domain, GRID, T-cell;
co-stimulatory adaptor protein; tissue rejection; graft rejection;
leukaemia; cytostatic; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New nucleic acid(s) for regulating the Grb2-related with Insert I (GRID) gene comprises using antisense and enzymatic nucleic acid molecules such as hammerhead ribozymes.
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                                                                                                                                                                                                                                                                                                                                                          Human GRID hammerhead ribozyme substrate oligonucleotide #75
                                                                              0.8%; Score 14; DB 1; Length 17;
100.0%; Pred. No. 8.3e+02;
ative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0.8%; Score 14; DB 1; Length 17; 100.0%; Pred. No. 8.3e+02; tive 0; Mismatches 0; Indels
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                                               Sequence 17 BP; 5 A; 5 C; 2 G; 0 T; 5 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mcswiggen JA,
                used to illustrate the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 4; Page 60; 108pp; English
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                                                                                                                                            598 TTTGGGAAACTGGA 611
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ABL46442 standard; RNA; 17
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                                                                   Query Match
Best Local Similarity 100.C
Marches 14; Conservative
                                                                                                                                                                                                                                                                                                                         27-JUN-2003 (first entry)
                                                                                                                                                                             TTTGGGAAACTGGA 2
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Best Local Similarity 100.
Matches 14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2001-550088/61.
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This invention describes a novel isolated nucleic acid that encodes one of three new isoforms of human pregnancy associated plasma protein E, hPAPP-E. The products of the invention have abortive and contraceptive activity and can be used for gene therapy or in a vaccine. The nucleic acid, polypeptide encoded by it, or antibody to the polypeptide can be used in pharmaceutical compositions or vaccines for preventing or aborting pregnancy. PAPP-E is used in the antenatal diagnosis of dysgenetic pregnancies. The nucleic acids are used as probes to assess the level of PAPP-E isoform mRNA in chorionic villus samples, and the antibodies can be used to assess the expression levels of PAPP-E isoform proteins in chorionic villus samples, to diagnose dysgenetic pregnancies antenatally. This sequence reperseents an oligomer used in scanning the human PAPP-E genes described in the disclosure of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PAPP-E; human; pregnancy associated plasma protein E; abortive; contraceptive; gene therapy; vaccine; pregnancy; antenatal; diagnosis; dysgenetic pregnancy; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                       New isolated nucleic acid encoding an isoform of human pregnancy associated plasma protein E, for preventing or aborting pregnancy
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 14; DB 1; Length 17;
Pred. No. 8.3e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        100.0%; Pred. ...
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human PAPP-Ea associated 17-mer SEQ ID 542.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 2; Page 146; 353pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ABS75016 standard; DNA; 17 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0.88;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        287 AACTICGITCIGCA 300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       4 AACTICGITCIGCA 17
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Best Local S
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Gaps

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Indels

598 TTTGGGAAACTGGA 611

TTTGGGAAACTGGA 1

g 8

RESULT 1242 ABS75015

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Homo sapiens

01-AUG-2002

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Gu Y,

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The invention relates to genetically modified fungus with an increased ability to utilise L-arabinose, where the fungus has been transformed with a DNA sequence encoding an L-arabinito1 4-dehydrogenase (EC 1.1. 1.12) or L-xylulose reductase (EC 1.1.1.10) or both the DNA sequences. Genetically modified fungus is useful for producing useful products from biomass containing L-arabinose. The useful product include ethanol, lactic acid or xylitol preferably ethanol. It is also useful to ferment a carbon source such as biomass comprising agricultural or forestry products and waste products containing L-arabinose and also other pentoses or other fermentable sugars. The present sequence is a PCR primer used to clone T. resesi L-arabinitol 4-dehydrogenase gene. (Updated on 29-AUG-2003 to standardise OS field)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Cytostatic; virucide; neuroprotective; nootropic; neuroleptic; gene chip; antisense; sense; tumour; cell degeneration; cancer; Alzheimer's disease; schizophrenia; protein chip; gene therapy; tumour suppression;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New isolated nucleic acid, useful for treating viral diseases associated with tumors and cell degeneration, also related polypeptides, antibodies and transfected cells.
                                                                                                                              Genetically modified fungus for producing useful products such as ethanol, lactic acid and xylitol, from biomass containing L-arabinose, has increased ability to utilize L-arabinose.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Tumour suppression related human fukutin oligo SEQ ID No 1839.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0.8%; Score 14; DB 1; Length 17;
100.0%; Pred. No. 8.3e+02;
ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 17 BP; 5 A; 1 C; 8 G; 3 T; 0 U; 0 Other;
(VALW ) VALTION TERNILLINEN TUTKIMUSKESKUS.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Tuijnder M;
                                                                                                                                                                                                                                   Example 2; Page 14; 32pp; English.
                                               Penttilae M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ABT36202 standard; DNA; 17 BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            17-SEP-2002; 2002WO-IB004208.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AGCGTAAAGGATGG 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    8 AGCGTAAAGGATGG 21
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Best Local Similarity 10...
Local 14; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2003-313353/30.
                                                                                           WPI; 2002-691618/74.
                                                  Londesborough J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WO2003025175-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              human fukutin;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        This invention describes a novel isolated nucleic acid that encodes one of three new isoforms of human pregnancy associated plasma protein E, hpapp-E. The products of the invention have abortive and contraceptive activity and can be used for gene therapy or in a vaccine. The nucleic acid, polypeptide encoded by it, or antibody to the polypeptide can be used in pharmaceutical compositions or vaccines for preventing or aborting pregnancy. Papp-E is used in the antenatal diagnosis of dysgenetic pregnancies. The nucleic acids are used as probes to assess the level of PAPP-E isoform mRNA in chorionic villus samples, and the proteins in chorionic villus samples, to diagnose dysgenetic pregnancies proteins in chorionic villus samples, to diagnose dysgenetic pregnancies antenatally. This sequence represents an oligomer used in scanning the human PAPP-E genes described in the disclosure of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Genetically modified fungus; L-arabinose; L-arabinitol 4-dehydrogenase; BC 1.1.1.12; L-xylulose reductase; BC 1.1.1.10; agricultural product; biomass; lactic acid; xylitol; forestry product; fermentable sugar; ethanol; enzyme; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         3900 PCR primer, to clone T. reesei L-arabinitol 4-dehydrogenase gene.
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                                                                                                                                                                                                                                                                                                                                                                                                    New isolated nucleic acid encoding an isoform of human pregnancy associated plasma protein E, for preventing or aborting pregnancy.
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                                                                                                                                                    06-APR-2001; 2001US-00827998
                                                                                                                                                                                                  26-MAY-2000; 2000US-0207456P
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Best Local Similarity 100.0
Matches 14; Conservative
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                                                                                                                                                                                                                                                                SHAN/) SHANNON M E.
                                                                                                                                                                                                                                                                                                                                                           WPI; 2002-697817/75.
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                                                                                                                                                                                                                                                                                                                     Shannon ME;
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                                                           US2002102252-A1
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29-AUG-2003 27-DEC-2002

AAD46160;

1244

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AAD46160

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Gaps ; 0

The invention tradects to a mover isolated 1, mer incident atto sequence, given in the specification, a sequence containing at least 15 consecutive nucleotides from the 17 mer sequence, a sequence, a sequence, a sequence that considering them t, at least 80 % identify to the 17 mer sequence, a sequence that tybridizes to them under highly stringent conditions, or the complement of any of them, or the corresponding RNA. The novel isolated mucleic caids of the invention are useful as probes and primers for detecting, identifying admitifying and/or amplifying a nucleic acid, e.g. as one component of a gene chip, in vitro as (anti) sense reagents, and for production of recombinant polypeptides. Any of the nucleic acids, containing the nucleic acids, cells containing the cortor or antibodies directed against the polypeptides are useful for preparation of pharmaceuticals for prevention and/or treatment of viral degeneration, specifically cancer but also Alzheimer's disease and containing the expression of the 17 mer nucleic acids in diseases. The polypeptides can also be used to generate antibodies, and contained samples is useful for disgnosis and/or prognosis of these contained contains the polypeptides and antibodies are useful as components of protein chips. The nucleic acid sequences of the invention can be used in gene there are therefore and antibodies are useful as components of protein chips. The nucleic acid sequences of the invention can be used in gene therapy. This polypucteetide sequence represents a tumour suppression contains and protein contains and antibodies are useful and antibodies. The invention relates to a novel isolated 17 mer nucleic acid sequence, \$

Sequence 17 BP; 4 A; 6 C; 6 G; 1 T; 0 U; 0 Other;

Score 14; DB 1; Length 17; Pred. No. 8.3e+02; 0; Mismatches 0; Indels 100.08; LL 1573 TCAGGCAGGCCAGC 1586 3 TCAGGCAGGCCAGC 16 Local Similarity 100.0 Query Match Best Loca Matches ð 셤

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Gaps

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ACA06338 standard; RNA; 17 BP 03-JUN-2003 (first entry) ACA06338; RESULT 1246 ACA06338

NFKB sub-unit modulating inozyme substrate #157.

Enzymatic nucleic acid; nuclear factor kappa B; NFKB; inozyme; zinzyme; G-cleaver; amberzyme; cancer; REL-A activity; breast cancer; human; lung cancer; prostate cancer; colorectal cancer; brain cancer; costate cancer; brain cancer; costate cancer; costate cancer; parcreatic cancer; cervical cancer; stomach cancer; brain cancer; melanoma; cervical cancer; head and neck cancer; ovarian cancer; melanoma; chemotherapy; paclitaxel; docetaxel; cisplatin; methotrexate; chemotherapy; paclitaxel; docetaxel; cisplatin; methotrexate; ovclophosphamide; doxorubin; fluorouracil carboplatin; edatrexate; ovclophosphamide; doxorubin; fluorouracil carboplatin; edatrexate; theumatoid arthritis; restenosis; foron; selesses; obsetty; ischaemia; gene therapy; autoimmune disease; lupus; multiple sclerosis; sepsis; transplant/graft rejection; reperfusion injury; glomerulonephritis; allergic airway inflammation; inflammatory bowel disease; infection; ss.

Homo sapiens

US2002177568-A1.

28-NOV-2002

23-MAY-2001; 2001US-00864785.

94US-00245466. 94US-00291932. 96US-00777916. 92US-00987132. 07-DEC-1992; 18-MAY-1994; 15-AUG-1994; 23-DEC-1996;

(STIN/) STINCHCOMB D T. (MCSW/) MCSWIGGEN J. (/NILS)

(DRAP/) DRAPER K G.

Draper KG Mcswiggen J, Stinchcomb DT,

WPI; 2003-340953/32.

Novel enzymatic nucleic acid molecules which down regulates expression of a sequence encoding a subunit of nuclear factor kappa B useful for treating cancer, inflammatory disorders and autoimmune diseases.

Claim 3; Page 29; 72pp; English.

The invention describes an enzymatic nucleic acid molecule (I) which down regulates expression of a sequence encoding a subunit of nuclear factor Kappa B (NFRB), where (I) is an incayme, zinzyme, G-cleaver or amberzyme configuration. The enzymatic nucleic acid molecule is adapted to treat cancer and is useful for down-regulating REL-A activity in a cell, for treating a patient having a condition associated with the level of REL-A. (I) is useful for cleaving RNA comprising a sequence of REL-A gene, in the presence of a divalent cation, especially MG'2+. The enzymatic and autisense nucleic acid molecules are useful for treating breast, lung, prostate, colorectal, brain, oesophageal, stomach, bladder, pancreatic, cervical, head and neck, ovarian cancer, melanoma, lymphoma, glioma or therapies such as monoclonal antibodies, REL-A-specific inhibitors or hown-therapies such as monoclonal antibodies, REL-A-specific inhibitors or chemotherapy including pacitizate, docetaxel, cigplatin, methotrexate, cyclophosphamide, doxorubin, fluorouracil carboplatin, edatrexate, gemcitabine or radiation therapy. The enzymatic and antisense nucleic acid molecules are also useful for treating inflammatory disease such as rheumatoid arthritis, restenosis, asthma, Crohn's disease, diabetes, rejection, gene therapy applications, ischaemia/reperfusion injury rejection, gene therapy applications, ischaemia/reperfusion injury (central nervous system (NS) and myocardial), glomerulonephritis, sepsis, allergic airway inflammation, inflammatory bowel disease or infection. This sequence represents the substrate of a novel enzymatic nucleic acid molecule

Sequence 17 BP; 5 A; 5 C; 1 G; 0 T; 6 U; 0 Other;

.. 0 0.8%; Score 14; DB 1; Length 17; 71.4%; Pred. No. 8.3e+02; tive 4; Mismatches 0; Indels 10; Conservative Local Similarity Query Match Matches

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Gaps

538 CCCATCTTTGACAA 551 3 CCCAUCUUGACAA 16 g

Human H-Ras DNAzyme target #115. ABZ61324 standard; RNA; 17 (first entry) 21-MAR-2003 ABZ61324; RESULT 1247 ABZ61324

BP

Human; ribozyme; short interfering RNA; siRNA; HER2; K-Ras; enzymatic nucleic acid; H-Ras; N-Ras; HIV; cytostatic; anti-HIV; anti-rheumatic; cancer; AIDS; ss.

Homo sapiens

WO200297114-A2.

05-DEC-2002.

29-MAY-2002; 2002WO-US016840.

29-MAY-2001; 2001US-0294140P. 06-JUN-2001; 2001US-0296249P. 10-SEP-2001; 2001US-0318471P.

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Claim
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 * X D D D D D D D D D D D X X X
                                                                                                                                             The invention relates to a novel short interfering RNA (siRNA) nucleic acid molecule or an enzymatic nucleic acid molecule, that modulates expression of a nucleic acid molecule encoding HRS2, K-Ras, H-Ras, N-Ras, human immunodeficiency virus (HIV) or a component of HIV. The nucleic acid molecule of the invention has cytostatic, anti-HIV, and anti-rheumatic activity. The nucleic acid molecules are useful for reducing HER2, K-Ras, H-Ras, and HIV activity in a cell. The nucleic acids are also useful for treating breast, ovarian, colorectal, lung, prostate, bladder, or pancreatic cancer, and HIV infection, and AIDS. The sequences shown in ABZ59889 - ABZ62216, ABZ6551, ABZ65520 - ABZ65524, ABZ66530 - ABZ6658 represent substrate/target sequences for the human ribozymes of the invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Novel short interfering RNA and enzymatic nucleic acid useful for treating cancer, modulates the expression of a nucleic acid encoding HER2, K-Ras, H-Ras, N-Ras, and human deficiency virus sequences.
                                                                                                                                                                                                                                                                                                                                                 Gaps
                                                                                Novel short interfering RNA and enzymatic nucleic acid useful for treating cancer, modulates the expression of a nucleic acid encoding HER2, K-Ras, H-Ras, N-Ras, and human deficiency virus sequences.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human; ribozyme; short interfering RNA; siRNA; HER2; K-Ras; enzymatic nucleic acid; H-Ras; N-Ras; HIV; cytostatic; anti-HIV; anti-rheumatic; cancer; AIDS; ss.
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                                                                                                                                                                                                                                                                                                                          0.8%; Score 14; DB 1; Length 17;
100.0%; Pred. No. 8.3e+02;
tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                      Sequence 17 BP; 2 A; 11 C; 4 G; 0 T; 0 U; 0 Other;
                                                                                                                               Claim 58; Page 113; 185pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human H-Ras DNAzyme target #970.
                                                                                                                                                                                                                                                                                                                                                                                                                                                         BP.
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06-JUN-2001; 2001US-0296249P.
10-SEP-2001; 2001US-0318471P.
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                                                                                                                                                                                                                                                                                                                                                                          103 CGCGCGCCCCGGC 116
                                                                                                                                                                                                                                                                                                                                                                                                                                                         ABZ62179 standard; RNA; 17
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                (RIBO-) RIBOZYME PHARM INC
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                                                             WPI; 2003-140484/13
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mcswiggen J;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           05-DEC-2002
                                       Mcswiggen J;
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                                                                          The invention relates to a novel short interfering RNA (siRNA) nucleic acid molecule or an enzymatic nucleic acid molecule, that modulates expression of a nucleic acid molecule encoding HER2, K-Ras, H-Ras, N-Ras, human immunodeficiency virus (HIV) or a component of HIV. The nucleic acid molecule of the invention has cytostatic, anti-HIV, and anti-rheumatic activity. The nucleic acid molecules are useful for reducing HER2, K-Ras, H-Ras, and HIV activity in a cell. The nucleic acids are also useful for treating breast, ovarian, colorectal, lung, prostate, bladder, or pancreatic cancer, and HIV infection, and AIDS. The sequences shown in ABZ59889 - ABZ62216, ABZ65531, ABZ65530 - ABZ66524, ABZ66530 - ABZ66530 - ABZ66524, ABZ66530 - ABZ66530 - ABZ66524, ABZ66531, ABZ66530 - ABZ66524, ABZ66530 - ABZ66531, ABZ66530 - ABZ66524, ABZ66530 - ABZ66530 - ABZ66534, ABZ66530 - ABZ66534, ABZ66530 - ABZ66534, ABZ66530 - ABZ66534, ABZ66534, ABZ66530 - ABZ66534, ABZ66534
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        p450,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Cancer; CYP3A5; irinotecan; pharmaceutical; malignant glioma; cytochrome p450; subfamily IIIA; nifedipine oxidase; polypeptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0.8%; Score 14; DB 1; Length 17;
100.0%; Pred. No. 8.3e+02;
tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 17 BP; 2 A; 5 C; 4 G; 0 T; 6 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (EPID-) EPIDAUROS BIOTECHNOLOGIE AG.
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58; Page 131; 185pp; English.
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2002EP-00011710.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ACF62527 standard; DNA; 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ribozymes of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        cytostatic; PCR primer; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
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Best Local Similarity
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24-MAY-2002;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  20-FEB-2003.
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52 GCAGTGTGACTGCTGA

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potentially harmful treatment of those subjects who do not respond to the treatment with substances (nonresponders), as well as the development of drug resistances due to suboptimal drug dosing can be avoided. ACF62200 to ACF62751 and ABM34912 to ABM35013 represent sequences used in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 or
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           irinotecan, colorectal cancer; cervical cancer; gastric cancer;
lung cancer; ovarian cancer; pancreatic cancer; malignant glioma;
variant allele; multidrug resistance protein 1; MRP1; cytostatic; gene;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Use of irinotecan or its derivative for preparation of a pharmaceutical composition for treating cancer in a subject having a genome with a variant allele comprising a multidrug resistance protein 1
                                                                                                                                                                                                                                                                                                                                                                      Gaps
                                                                                                                                                                                                                                                                                                                                                                   0;
                                                                                                                                                                                                                                                                                             Length 17;
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                                                                                                                                                                                                                Sequence 17 BP; 4 A; 3 C; 4 G; 5 T; 0 U; 1 Other;
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                                                                                                                                                                                                                                                                                      0.8%; Score 14; DB 1; I
87.5%; Pred. No. 8.3e+02;
live 1; Mismatches 1;
                                                                                                                                                  exemplification of the present invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADB21198 standard; DNA; 17 BP
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24-MAY-2002; 2002EP-00011710.
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                                                                                                                                                                                                                                                                                                                                                                                                                                 52 GCAGTGTGACTGCTGA
                                                                                                                                                                                                                                                                                                                   Local Similarity 87.5
les 14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2003-354397/33.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Unidentified.
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                                                                                                                                                                                                                                                                                          Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADB21198
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   8888888
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The invention relates to the novel use of irinotecan to treat a patient suffering from cancer. This involves determining if the patient has one or or more variant alleles of the UGTIAI gene, and if the patient has one or more of such variant alleles, irinotecan is administered in an increased or decreased amount in comparison to the amount that is administered or decreased amount in comparison to the amount that is administered without regard to the patient's alleles in the UGTIAI gene. The invention has cytostatic activity. A composition of the invention acts as a topologomerase I inhibitor. The method is useful for treating a patient, an animal e.g. mouse or a human, preferably African or Asian, suffering
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   from cancer such as colorectal, cervical, gastric cancer, lung, ovarian, pancreatic cancer or malignant glioma. The present sequence is udes in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ss; irinotecan; cancer; UGT1A1; cytostatic; topoisomerase I inhibitor; colorectal cancer; dervical cancer; gastric cancer; lung cancer; ovarian cancer; pancreatic cancer; malignant glloma; uridine diphosphate glycosyltransferasel member Al.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
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0
                                                                                                                                                                                                                                                                                                                                                                                                                                 Human UGT1A1 variant allele sequence fragment SEQ ID NO:328.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (EPID-) EPIDAUROS BIOTECHNOLOGIE AG.
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                                                                                                                                                                                                                    BP
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2 GCAATGTRACTGCTGA 17
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24-MAY-2002; 2002EP-00011710.
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                                                                                                                                                                                                         ADB88287 standard; DNA; 17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           14; Conservative
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ADB97270;

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Gaps

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0.8%; Score 14; DB 1; Length 17; 87.5%; Pred. No. 8.3e+02; tive 1; Mismatches 1; Indels

Query Match 0.8 Best Local Similarity 87.5 Matches 14; Conservative

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New use of irinotecan for preparation of pharmaceutical compositions for treating cancer in subject having genome with variant allele comprising multidrug resistance 1 polynucleotide.
                                                                                                                                                                                                                                                                                                                                                                                                        The invention relates ro a novel use of irinotecan or its derivative for the preparation of a pharmaceutical composition for treating colorectal,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human; ss; GRID; Grb2-related with insert domain; hammerhead ribozyme; NCH ribozyme; G-cleaver ribozyme; Zinzyme; DNAzyme; amberzyme; Inozyme; hairpin ribozyme; tissue rejection; graft rejection; leukaemia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                    cervical, gastric, lung, ovarian or pancreatic cancer, or malignant glioma in a subject having a genome with a variant allele which comprise multidrug resistance 1 (MDR1) polynucleotide. A composition of the invention has cytostatic activity. The present sequence is used in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0.8%; Score 14; DB 1; Length 17;
llarity 87.5%; Pred. No. 8.3e+02;
Conservative 1; Mismatches 1; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human GRID mRNA substrate sequence #73.
                                                                                                                                                                                                                                                                                                                                                                          Disclosure, Page 50; 104pp; English.
                                                                                                                                                                (EPID-) EPIDAUROS BIOTECHNOLOGIE AG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             exemplification of the invention.
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                                                        23-JUL-2002; 2002WO-EP008220.
                                                                                                 23-JUL-2001; 2001EP-00117608.
24-MAY-2002; 2002EP-00011710.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                52 GCAGTGTGACTGCTGA 67
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(CARL/) CARLOWITZ I V.
(MCSW/) MCSWIGGEN J.
(HAMB/) HAMBLIN P A.
(ELLI/) ELLIS J H.
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nes 14; Conserv
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                   20-FEB-2003.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The invention relates to the novel use of irinotecan or its derivative for the preparation of pharmaceutical compositions for treating colorectal, cervical, gastric, lung, ovarian or pancreatic cancer, or malignant glioma in a subject having a genome with a variant allele which comprises a multidrug resistence 1 (MDR1) polynuclectide. A composition of the invention has cytostatic activity. The invention is useful for the preparation of pharmaceutical compositions for treating colorectal, cervical, gastric, lung, ovarian or pancreatic cancer, or malignant glioma in a subject (preferably human, more preferably African or Asian) or a mouse. The present sequence is used in the exemplification of the
                                                                                                   irinotecan; colorectal cancer; cervical cancer; gastric cancer; lung cancer; ovarian cancer; pancreatic cancer; malignant glioma; multidrug resistance 1; MDR1; cytostatic; human; ds; Cyp3A5; MRP1; MDR1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New use of irinotecan for preparation of pharmaceutical compositions for treating cancer in subject having genome with variant allele comprising multidrug resistance 1 polynucleotide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      irinotecan; colorectal cancer; cervical cancer; gastric cancer;
lung cancer; ovarian cancer; pancreatic cancer; malignant glioma;
multidrug resistance 1; MDR1; cytostatic; ds; human; UGT1A1; MRP1; TOP1.
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                                                                 Human MDR1 variant allele sequence fragment SEQ ID NO:356.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0.8%; Score 14; DB 1; Length 17; llarity 87.5%; Pred. No. 8.3e+02; Conservative 1; Mismatches 1; Indels
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24-MAY-2002; 2002EP-00011710
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                          (first entry)
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Best Local Similarity
Matches 14; Conserv
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Gaps . 0

Ellis JH;

10017621-3sl.rng

New nucleic acid molecule that down-regulates expression of Grb2-related with insert domain (GRID) gene, useful for treating a condition associated with the level of GRID, e.g. tissue/draft rejection and gene, useful for treating a condition of GRID, e.g. tissue/graft rejection and

Claim 4; SEQ ID NO 73; 74pp; English.

The invention relates to a nucleic acid molecule that down-regulates expression of Grb2-related with insert domain (GRLD) gene, e.g. a hammerhead ribozyme, NCH ribozyme, G-cleaver ribozyme, Zinzyme, DNAzyme, amberzyme, Inozyme or hairpin ribozyme. Also include are a mammalian cell including the novel nucleic acid molecule, reducing GRID activity in a cell by contacting the cell with the novel nucleic acid molecule, treathing a patient having a condition associated with the level of GRID (e.g. tissue/graft rejection or leukaemia) by contacting the cell with the novel nucleic acid molecule, cleaving RNA of a GRID gene by contacting the cell with the novel nucleic acid molecule, an expression vector comprising a nucleic acid sequences (encoding at less the novel nucleic acid molecule in a manner that allows its expression), a mammalian cell including the expression vector and an enzymatic nucleic acid molecule that cleaves RNA derived from a GRID gene. The nucleic acid molecule is useful for treating a condition associated with the level of GRID, e.g. tissue/graft rejection and leukaemia. The present sequence is a target region for the enzymatic nucleic acids of the invention.

Sequence 17 BP; 5 A; 6 C; 1 G; 0 T; 5 U; 0 Other;

Gaps ·, 0.8%; Score 14; DB 1; Length 17; 100.0%; Pred. No. 8.3e+02; tive 0; Mismatches 0; Indels Best Local Similarity 100. Matches 14; Conservative Query Match ઠે

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598 TTTGGGAAACTGGA 611 TTTGGGAAACTGGA 3

Db

ADM53799/c

ADM53799 standard; mRNA; 17 BP ADM53799 HANDER STANDER STANDER

(first entry) 03-JUN-2004

Human GRID mRNA substrate sequence #74.

Human, ss, GRID, Grb2-related with insert domain; hammerhead ribozyme; NCH ribozyme; G-cleaver ribozyme; Zinzyme; DNAzyme; amberzyme; Inozyme; hairpin ribozyme; tissue rejection; graft rejection; leukaemia.

Homo sapiens

US2003134806-A1.

17-JUL-2003

23-FEB-2001; 2001US-00792818.

10-FEB-2000; 2000US-0181594P.

JARVIS T.
CARLOWITZ I V.
MCSWIGGEN J. CARL/) (MCSM/)

JARV/)

HAMBLIN P A. ELLIS J H. (ELLI/) (HAMB/)

Mcswiggen J, Carlowitz IV, Jarvis T,

Ellis JH;

Hamblin PA,

WPI; 2003-829646/77.

New nucleic acid molecule that down-regulates expression of Grb2-related with insert domain (GRID) gene, useful for treating a condition associated with the level of GRID, e.g. tissue/graft rejection and

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The invention relates to a nucleic acid molecule that down-regulates expression of Grb2-related with insert domain (GRID) gene, e.g. a harmeribad ribozyme, NGH ribozyme, G-cleaver ribozyme, Zinzyme, DNAzyme, amberzyme, Inozyme or hairpin ribozyme. Also include are a mammalian cell including the novel nucleic acid molecule, reducing GRID activity in a cell by contacting the cell with the novel nucleic acid molecule, reducing GRID activity in a treating a patient having a condition associated with the level of GRID (e.g. tissue/graft rejection or lenkaemia) by contacting the cell with the novel nucleic acid molecule, an expression contacting the cell with the novel nucleic acid molecule, an expression vector comprising a nucleic acid sequences (encoding at least the novel nucleic acid molecule in a manner that allows its expression), a mammalian cell including the expression vector and an enzymatic nucleic acid molecule that cleaves RNA derived from a GRID gene. The nucleic acid molecule that cleaves RNA derived from a GRID gene. The nucleic acid molecule that cleaves RNA derived from a GRID gene. The nucleic acid molecule is useful for treating a condition associated with the level of GRID, a traver ranion for the contaction and leukaemia. The present sequence is

    tissue/graft rejection and leukaemia. The present sequence is
region for the enzymatic nucleic acids of the invention.

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100.0%; Pred. No. 8.3e+02;
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                                                  Claim 4; SEQ ID NO 74; 74pp; English.
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leukemia.
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ADM53800 standard; mRNA; 17 BP (first entry) 03-JUN-2004 ADM53800; RESULT 1256 ADM53800/c

Human GRID mRNA substrate sequence #75.

Human, 88, GRID, Grb2-related with insert domain, hammerhead ribozyme, NCH ribozyme, G-cleaver ribozyme, Zinzyme, DNAzyme, amberzyme, Inozyme, hairpin ribozyme, tissue rejection, graft rejection, leukaemia.

Homo sapiens

US2003134806-A1.

17-JUL-2003.

23-FEB-2001; 2001US-00792818. 10-FEB-2000; 2000US-0181594P.

CARLOWITZ I V. MCSWIGGEN J. HAMBLIN P A. ELLIS J H. JARV/) JARVIS T. (ELLI/) CARL/) MCSM/) (HAMB/

Ellis JH; Hamblin PA, Mcswiggen J, Carlowitz IV, Jarvis T,

WPI; 2003-829646/77.

New nucleic acid molecule that down-regulates expression of Grb2-related with insert domain (GRID) gene, useful for treating a condition associated with the level of GRID, e.g. tissue/graft rejection and Leukemia.

Claim 4; SEQ ID NO 75; 74pp; English

Sequence 17 BP; 4 A; 5 C; 3 G; 0 T; 5 U; 0 Other;

The invention relates to a nucleic acid molecule that down-regulates expression of GrD2-related with insert domain (GRID) gene, e.g. a hammerhead ribozyme, NCH ribozyme, G-cleaver ribozyme, Zinzyme, DNDxyme, amberzyme, Inozyme or hairpin ribozyme. Also include are a mammalian cell including the novel nucleic acid molecule, reducing GRID activity in a cell by contacting the cell with the novel nucleic acid molecule, treating a patient having a condition associated with the level of GRID (e.g. tissue/graft rejection or leukaemia) by contacting the cell with the novel nucleic acid molecule, cleaving RNA of a GRID gene by contacting the cell with the novel nucleic acid molecule, acid molecule, an expression vector comprising a nucleic acid sequences (encoding at least the novel nucleic acid molecule in a manner that allows its expression), a mammalian cell including the expression vector and an enzymatic nucleic acid molecule that cleaves RNA derived from a CRID gene. The nucleic acid molecule is useful for treating a condition associated with the level of GRID, e.g. tissue/graft rejection and leukaemia. The present sequence is a target region for the enzymatic nucleic acids of the invention.

Score 14; DB 1; Length 17; 8.3e+02; hes 0; 100.0%; Pred. No. 8.3 tive 0; Mismatches 0.8%; 14; Conservative Query Match Best Local Similarity Matches

598 TITGGGAAACTGGA 611

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Gaps .; 0

Indels

14 TTTGGGAAACTGGA 1 음

Human KDR VEGF receptor hairpin ribozyme substrate #40. AAX71742 standard; RNA; 18 BP. (first entry) 28-JUL-1999 AAX71742; RESULT 1257

Vascular endothelial growth factor receptor; VEGF receptor; flt-1; flk-1; KDR; hammerhead ribozyme; hairpin ribozyme; cleavage; tumour angiogenesis; psoriasis; rheumatoid arthritis; ocular disease; fms-like tyrosine kinase 1; kinase insert domain containing receptor; foetal liver kinase 1; ss.

Homo sapiens

WO9715662-A2

01-MAY-1997,

96WO-US017480 25-OCT-1996; 95US-0005974P. 26-0CT-1995; 11-JAN-1996;

(RIBO-) RIBOZYME PHARM INC CHIRON CORP (CHIR) Escobedo J; Ď, Stinchcomb Mcswiggen J, Pavco P,

WPI; 1997-259017/23.

Nucleic acid molecule modulating VEGF receptor(s) gene expression or mRNA stability - useful for treating e.g. tumour angiogenesis, psoriasis, rheumatoid arthritis, etc., in a human patient.

Claim 4; Page 120; 218pp; English.

The present invention describes nucleic acid molecules which modulate the synthesis, expression and/or stability of a mRNA encoding 1 or more receptors of vascular endothelial growth factor (VEGF). A patient

A method has been developed of defining a set of compounds that modulate the expression of a target nucleic acid (tNA) sequence via binding of the compounds with the tNA sequence. The method comprises generating a evaluating in silico the binding of the virtual compounds with the tNA according to defined criteria. Also described are: (1) a method of according to defined criteria. Also described are: (1) a method of a defining a set of oligonucleotides (ONS) that modulate the expression of a tNA sequence via binding of the ONS with the tNA sequence comprising generating a library of virtual compounds in silico according to defined criteria, and evaluating in silico the binding of the virtual ONS with the tNA according to defined criteria; and (2) a method of defining a set of compounds that modulate the expression of a tNA sequence via binding of the compounds with the tNA. The methods can be used for the generation and identification of synthetic compounds having defined physical, ö to рe treated by administering the nucleic acid molecule or the expression vector to the patient. AAX67275 to AAX75752 represent specific examples of nucleic acid molecules from the present invention nsed fms-like tyrosine kinase I (flt-1), kinase insert domain containing receptor (KDR) and/or foetal liver kinase 1 (flk-1) (e.g. tumour angiogenesis, ocular diseases, psoriasis and rheumatoid arthritis) can Brooks DG; Human ELK-1 phosphorothioate antisense oligonucleotide SEQ ID NO:206. Gaps preferably human) having a condition associated with the level of Identifying compounds which modulate expression of nucleic acids, provide compounds having defined physical, chemical or bioactive properties, e.g. antisense activity. Identification; genetic target; gene modulation; human; probe; antisense oligonuclectide; phosphorothicate; PCR primer; nucleotide sequence-based technology; antisense drug discovery; .. Sasmor HM, Score 14; DB 1; Length 18; Pred. No. 8.7e+02; 0; Indels Sequence 18 BP; 2 A; 9 C; 1 G; 0 T; 6 U; 0 Other; Freier SM, Vickers TA; 5; Mismatches Example 24; Page 104; 264pp; English. f, Mcneil J, Borchers AH, BP 98US-0081483P. 0.8%; 99WO-US008268 98US-00067638 CTCTCTGCCTACCT 1714 AAZ41054 standard; DNA; 18 cucucucccuaccu 15 (first entry) 9; Conservative (ISIS-) ISIS PHARM INC. BF, target validation; ss. Cowsert LM, Baker B Ohasi C, Wyatt JR, WPI; 1999-620446/53 Local Similarity 26-JAN-2000 Homo sapiens. W09953101-A1 13-APR-1999; L3-APR-1998; 28-APR-1998; 21-OCT-1999 Synthetic. AAZ41054; 1701 Query Match RESULT 1258 Matches 888888888 ò 원

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chemical or bioactive properties. Information gathered from assays of such compounds is used to identify nucleic acid sequences that are tractable to a variety of nucleotide sequence-based technologies, e.g. antisense drug discovery and target validation. AAZ40852 to AAX41220, and AAX52701 to AAX52706, represent sequences used in the exemplification of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human ELK-1; p62TCF; Ets domain transcription factor protein, apoptosis, expression inhibition; infection; inflammation; tumour formation, diagnosis; phosphorothioate; antisense compound; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /*tag= c
/note= "Optionally 2-methoxyethyl (2'-MOE) nucleosides
except cytosine residues which are 5-methylcytosine"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /note= "Optionally 2-methoxyethyl (2'-MOE) nucleosides
except cytosine residues which are 5-methylcytosine"
                                                                                                                                                                                                                                                                                                                                   Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1. .18
  **tag= a
/*tag= "Internucleoside phosphorothioate linkages"
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                                                                                                                                                                                                          Sequence 18 BP; 0 A; 2 C; 12 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    location/Qualifiers
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vative 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ELK-1 expression modulator #9
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                                                                                                                                                                                                                                                                       0.88:
                                                                                                                                                                                                                                                                                                                                                                                232 GGTGGTGGTGGCGG 245
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                                                                                                                                                                                                                                                                                                                                                                                                                                       1 derecreerecee 14
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                                                                                                                                                                                                                                                                                                                       14; Conservative
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                                                                                                                                                    invention
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                                                                                                                                                                                                                                                                                               Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAZ06571;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Baker BF,
                                                                                                                                                                                                                                                                    Query Match
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AAZ06571
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Sequences AAZ06571-Z06607 are antisense polynucleotides targeted to a nucleic acid molecule encoding human ELK-1 (also known as p62TCF). ELK-1 is a member of the ternary complex factor subfamily of Ets-domain transcription factor proteins. The polynucleotides inhibit the expression of human ELK-1, and this sequence targets the 5' untranslated region of the ELK-1 RNA. Sequences AAZ06571-Z06607 all cause at least 30% inhibition of ELK-1 expression. The antisense sequences can be used to

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inhibit the expression of human ELK-1 in human cells or tissues in vitro. BLK-1 uses a bipartite recognition mechanism mediated by both protein-DNA and protein-protein interactions to regulate genes by direct and indirect DNA binding and has been shown to control various signal transduction pathways and other cell functions including apoptosis. This means that antisense compounds inhibiting expression of ELK-1 can be used to treat diseases associated with its expression in animals, particularly humans and to prevent or delay infection, inflammation or tumour formation. The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    useful for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 This invention describes a novel method for determining the degree of methylation of a particular cytosine in a motif 5'-CpG-3', present in a genomic sample of DNA. The sample is treated chemically to convert cytosine (C) but not methylated C, to uracil, then part of the genomic DNA that contains the target C is amplified to form a labeled amplicon. The amplicon is hybridised to two classes, each with at least one member, of oligonucleotides and/or peptide-nucleic acid (DNA) oligomes and the degree of hybridisation to both classes is determined from the label on the amplicon. From the ratio of labels hybridised to the two classes of oligomers, the degree of methylation is calculated. The method is used: (i) for diagnosis and/or prognosis of side effects of therapeutic drugs and of a wide range of diseases, e.g. cancer, disorders of the central nervous, cardiovascular, gastrointestinal and respiratory systems etc.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Determining the degree of cytosine methylation in genomic DNA, useful for diagnosis and prognosis, comprises selective hybridization of amplicons
                                                                                                                                                                     compounds can also be used for diagnosis, as research reagents and in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human; cytosine methylation; 5'-CpG-3'; uracil; cytosine; diagnosis; drug; side effect; cancer; central nervous system; cardiovascular; gastrointestinal; respiratory system; single nucleotide polymorphism; SNP; cell differentiation; ELK-1; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                   Gaps
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                                                                                                                                                                                                                                                                      0.8%; Score 14; DB 1; Length 18; 100.0%; Pred. No. 8.7e+02; tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                Sequence 18 BP; 0 A; 2 C; 12 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Guetig D;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Berlin K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 1; Page 33; 56pp; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ABA99961 standard; DNA; 18 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      01-SEP-2001; 2001WO-EP010074.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            01-SEP-2000; 2000DE-01043826.
05-SEP-2000; 2000DE-01044543.
                                                                                                                                                                                                                                                                                                                                                     232 GGTGGTGGTGGCGG 245
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           from chemically treated DNA.
                                                                                                                                                                                                                                                                                                                                                                                           1 deregreeregeeg 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human ELK-1 PCR primer #2.
                                                                                                                                                                                                                                                                                       Local Similarity 100.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Ú
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2002-371829/40.
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                                                                                                                                                                                                                                                                      Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                      RESULT 1260
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This invention describes a novel method for detecting cytosine

methylation in DNA samples by: (i) chemically treating a genomic sample

to convert all non-methylated cytosines to uracil while leaving

methylated cytosines unchanged; (ii) amplification with 2 primer

oligonucleotides and a polymerase; and (iii) analysis of the amplicon and

determing the methylation status of test DNA. The method is used for

deducing the methylation status at different CpG positions, which is

cused for diagnosis and/or prognosis of a very wide range of disorders,

cused for diagnosis and/or prognosis of a very wide range of disorders,

cused for diagnosis and/or infocutions, inflammation, etc. The method is

cused for diagnosis and infections, inflammation, etc. The method is

also useful for differentiating between cell and tissue types and for

investigating cell differentiation. The methylated positions, and

cunvertiative indication of the different methylated positions, and

covery accurate classification; and (ii) eliminates interference from

the ackground DNA, making it suitable for analysis of serum or body fluids

covery accurate classification; and (ii) eliminates interference from

which contain background DNA in large excess). This sequence represents

ca PCR primer used to amplify the human ELK-I gene, described in the
                                                                                                                                                                                                                                         ·
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particularly by detecting mutations or single nucleotide polymorphisms (SNP's); and (ii) for differentiation of cell or tissue types and for investigating cell differentiation. The method allows the methylation status of many C residues to be determined simultaneously. This sequence represents a PCR primer used in the amplification of the human ELK-1 gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human, cytosine methylation, methylation status; CpG; infection; cancer; diagnosis; side-effect, cardiovascular disease; gastrointestinal disease; inflammation; cell differentiation; ELK-1; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Detecting methylation status of test DNA in a mixture, useful for diagnosis and prognosis of disease, comprises bisulfite treatment then selective amplification of test DNA.
                                                                                                                                                                                                                                           Gaps
                                                                                                                                                                                                                                           ;
                                                                                                                                                                                                 DB 1; Length 18;
8.7e+02;
                                                                                                                                                                                                                                           Indels
                                                                                                                                                    Sequence 18 BP; 1 A; 1 C; 12 G; 4 T; 0 U; 0 Other;
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                                                                                                                                                                                              / Match 0.8%; Score 14; DB Local Similarity 100.0%; Pred. No. 8.7 nes 14; Conservative 0; Mismatches
                                                                                                              used in the method of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example 4; Page 43; 82pp; German
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human ELK-1 PCR primer SEQ ID 2.
                                                                                                                                                                                                                                                                                                                                                                                                                                   BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                09-MAR-2001; 2001DE-01012515.
19-NOV-2001; 2001DE-01058283.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        08-MAR-2002; 2002WO-EP002572.
                                                                                                                                                                                                                                                                                       232 GGTGGTGGTGGCGG 245
                                                                                                                                                                                                                                                                                                                                                                                                                                     AAF88946 standard; DNA; 18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Berlin K;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAF88946;
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differentiating between lung cell proliferative disorders associated with at least one gene and/or their regulatory regions. Specifically, it refers to a method comprising contacting a target nucleic acid in a biological sample with at least one reagent, wherein the reagent is able to distinguish between methylated and non-methylated cpd dinucleotides present in the target DNA. As such, it is possible to further differentiate and diagnose medical conditions including adenocarcinoma and squamous cell carcinoma, and their respective adjacent lung tissue. The present invention describes cytostatic oligomers and RNA-oligomers that are useful as probes for determining the cytosine methylation state or single nucleotide polymorphisms (SNPS) of the target sequence. This cyclic positions within genomic DNA, used in an exemplification of the
                                                                          ö
                                                                                                                                                                                                                                                                                                                                                                                 squamous cell carcinoma; cytostatic; probe; PNA-oligomer;
                                                                                                                                                                                                                                                                                                                              Primer oligo used for analysing CpG islands in genomic DNA (SeqID 771).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Detecting and differentiating cytosine methylation state of genomic DNP nearing for diagnosing, treating brognosticating and/or monitoring lung
                                                                            Gaps
                                                                                                                                                                                                                                                                                                                                                              PCR; primer; ss; lung cell proliferative disorder; CpG dinucleotide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ll for diagnosing, treating prognosticating and or monitoring proliferative disorders e.g. adenocarcinoma and squamous cell
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Maier S;
                                                                            ..
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   This invention relates to a novel method for detecting and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 14; DB 1; Length 18;
Pred. No. 8.7e+02;
                                          Length 18;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 18 BP; 3 A; 0 C; 10 G; 5 T; 0 U; 0 Other;
        Sequence 18 BP; 1 A; 1 C; 12 G; 4 T; 0 U; 0 Other;
                                          Score 14; DB 1; L. Pred. No. 8.7e+02;
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                                                                             Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 15; SEQ ID NO 771; 58pp; English.
                                            0.8°,
100.0%; Pre-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Genc B,
                                                                                                                                                                                                                                    BP.
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                                                                                                                 GGTGGTGGTGGCGG 245
                                                                                                                                                                                                                                                                                                                                                                                                     cytosine methylation state.
                                                                                                                                                                                                                                    ADC70281 standard; DNA; 18
                                                                                                                                               derecrecrededed 15
                                                            Local Similarity 100.
nes 14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (EPIG-) EPIGENOMICS AG.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Field JK,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2003-533029/50.
                                                                                                                                                                                                                                                                                                                                                                                       adenocarcinoma;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WO2003052135-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                          Unidentified.
                                                                                                                                                                                                                                                                                                     18-DEC-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           26-JUN-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   useful for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     carcinoma
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Burger M,
                                                                                                                                                  N
                                                                                                                                                                                                                                                                     ADC70281;
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                                                                                                                  232
                                               Query Match
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                                                                   Best Loca
Matches
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1156 ATGTGGGGTGTGGG 1169

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New polynucleotides encoding secreted human proteins, derived from human fetal brain, adult skin, adult brain, adult heart, adult thymus and adult aorta cDNA libraries.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         This invention describes novel human secreted proteins which are encoded by polynucleotides obtained from fetal brain, adult skin, adult brain, adult heart, adult thymus and adult neart constitutions. The polynucleotides and proteins are predicted to have biological activities which would make them suitable for treating, preventing or ameliorating medical conditions in humans and animals, although no supporting data is given. Suggested activities include nutritional activity, cytokine and cell proliferation/differentiation activity, immune stimulating (e.g. as vaccines) or suppressing activity, hematopoiesis regulating activity, chemotractic/chemokineic activity, hemostatic and thrombolytic activity, receptor/ligand activity, anti-inflammatory activity, cadherin/tumor activity, anti-inflammatory activity, cadherin/tumor activity, anti-inflammatory activity, the
                                                                                                                                                                                                                     Human, secreted protein, treatment, nutritional activity, cytokine, cell proliferation, cell differentiation, hematopoissis regulation, tissue growth, activin, inhibin, chemotactic, chemokinetic, hemostatic, thrombolytic, anti-inflammatory, invasion suppressor, tumor inhibition,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       polynuclectides are also stated to be useful for gene therapy. AAZ43809-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       243840 represent DNA probes used to isolate the polynucleotides represented in AAZ43777-Z43808 which encode the secreted proteins represented in AAX50905-Y50947
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 19 BP; 6 A; 1 C; 10 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                      Human adult thymus cDNA clone vhl_1 DNA probe.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Disclosure; Page 270; 282pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Hoffman H,
                                                                                     BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               98US-0082904P.
98US-008934P.
98US-0089278P.
98US-0091647P.
98US-0097639P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                 99WO-US008504
                                                                                  AAZ43839 standard; DNA; 19
 ATGTGGGGTGTGGG 14
                                                                                                                                                    10-MAR-2000 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Yuan O,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (ALPH-) ALPHAGENE INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2000-052801/04.
                                                                                                                                                                                                                                                                                            gene therapy; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Valenzuela D,
                                                                                                                                                                                                                                                                                                                                              Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                               WO9955721-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                 23-APR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   12-JUN-1998;
                                                                                                                                                                                                                                                                                                                                                                                                               04-NOV-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   24-APR-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       02-JUL-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     22-APR-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     11-JUN-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     24-AUG-1998
                                                                                                                                                                                                                                                                                                                            Synthetic
                                                                                                                    AAZ43839;
                                                                  AAZ43839
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The present invention relates to a hairpin or hammerhead ribozyme, designed to cleave RNA encoding a cyclin or cell-cycle dependent kinase other than cell-cycle dependent kinases CDKI, PCNA and Cyclin BI.
Representative examples of ribozyme recognition sites are given in AAA8415 to AAA86787. The ribozyme of the invention is useful for inhibiting restenosis by introduction of the ribozyme into cells. The ribozyme is resistant to endonuclease activity and hence is efficient in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human; ribozyme therapy, hairpin ribozyme; hammerhead ribozyme; recognition site; target; ribozyme binding site; eye disease; vulnerary; proliferative disease; skin disease; psoriasis; diabetic retinopathy; cytokine; inflammation; cell-cycle dependent kinase; cyclin; MVP; matrix metalloproteinase; growth factor; reductase; scarring; cytostatic;
                                                                                                                                   Ribozyme; hairpin; hammerhead; gene therapy; vasotropic; restenosis; ss.
                                                                                                                                                                                                                                                                                                                                                                                                     New hairpin and hammerhead ribozyme for inhibiting restenosis, cleaves RNA encoding a cyclin or cell-cycle dependent kinase other than CDK1, PCNA and Cyclin B1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Cell-cycle dependent kinase cdk2 ribozyme binding site SEQ ID NO:203.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0.8%; Score 14; DB 1; Length 19; 100.0%; Pred. No. 9.1e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 19 BP; 2 A; 6 C; 4 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                              Robbins JM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     100.0%; Prec. ...
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Disclosure; Page 49; 109pp; English.
                                                                                                                                                                                                                                                                                                                                                Barber JR,
                                                                                                    cdk2 ribozyme binding site #54
                AAA82617 standard; DNA; 19 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAH57779 standard; DNA; 19 BP.
                                                                                                                                                                                                                                                        99WO-US028772
                                                                                                                                                                                                                                                                                    98US-0110954P
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                                                                          (first entry)
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hes 14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CTGTTCCAGCTGCT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  restenosis treatment
                                                                                                                                                                                                                                                                                                                                           Tritz R, Welch PJ,
                                                                                                                                                                                                                                                                                                              (IMMU-) IMMUSOL INC.
                                                                                                                                                                                                                                                                                                                                                                         WPI; 2000-412314/35
                                                                                                                                                                                             WO200032765-A2
                                                                                                                                                                                                                                                       06-DEC-1999;
                                                                                                                                                                                                                                                                                  04-DEC-1998;
                                                                          04-DEC-2000
                                                                                                                                                                                                                           08-JUN-2000.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     10-SEP-2001
                                            AAA82617;
                                                                                                                                                                  Mammalia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      922
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAH57779;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             RESULT 1265
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Matches
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AAA8261
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Rapiejko P;

Hall J,

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Query Match
Best Local Similarity 100.
Matches 14; Conservative

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0.8%; Score 14; DB 1; Length 19; 100.0%; Pred. No. 9.1e+02;

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ADF37677 standard; RNA; 19 BP.
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                                                                                                                     WO2003070910-A2.
                                                                                             Homo sapiens.
                                                                                                                                                                                                                                                                                                                                     27-NOV-2002;
                                                                                                                                                                                                                                                                                                                                                                                                      Mcswiggen J,
                                                                                                                                                                                                    20-FEB-2002;
                                                                                                                                               28-AUG-2003
                                                                            Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local 9
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     d
                                                                                                                                                                                                                                                                                                                                                                                     The present invention describes a method for treating a proliferative skin or eye disease and scarring. The method involves administering a cribozyme (I) which cleaves RNA encoding a cytokine involved in inflammation, matrix metalloproteinase (MMP), cyclin, cell-cycle dependent kinase, growth factor or a reductase, or administering a nucleic acid molecule (II) comprising a promoter operably linked to a nucleic acid molecule (II) comprising a promoter operably linked to a characological, cytostatic, antiseborrheic, antidiabetic, antisickling, dermatological, vulnerary, keratolytic and virucide activities, and cleaves RNA encoding cycokine involved in inflammation. (I) can be used in gene therapy. (I) and (II) are useful for treating proliferative skin diseases such as psoriasis, atopic dermatitis, actinic keratosis, squamous or basal cell carcinoma and viral or seborrheic wart. They can also be used for treating proliferative eye diseases such as diabetic retinopathy, vitreoretinopathy, sickle cell retinopathy, retinopathy of carring and preventing and preventing and preventing carring and preventing methods.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ö
                                                                                                                                                                                                                                                                                                           Treating proliferative skin or eye diseases and scarring, using ribozymes that cleave RNA encoding cytokines involved in inflammation, matrix metalloproteinases, growth factors and cell-cycle dependent kinases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           or hypertrophic burn
antipsoriatic; dermatological; antiseborrheic; antidiabetic; virucide; antisickling; ophthalmological; keratolytic; gene therapy; viral wart; atopic dermatifis; actinic keratosis; squamous cell carcinoma; basal cell carcinoma; seborrheic wart; vitreoretinopathy; scar; sickle cell retinopathy; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             double-stranded short interfering nucleic acid;
short interfering nucleic acid; siNA; downregulation;
vascular endothelial growth factor receptor; VEGFR; antianglogenic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human VEGFR3 short interfering nucleic acid (siNA) SEQ ID NO:1719.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          sequences used in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0.8%; Score 14; DB 1; Length 19;
100.0%; Pred. No. 9.1e+02;
ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              scarring such as keloid, adhesion and hypertrophic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 19 BP; 2 A; 6 C; 4 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        exemplification of the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAH57577 to AAH62099 represent
                                                                                                                                                                                                                                                                                                                                                                   Example 1; Page 86; 408pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        100.08;
                                                                                                                                                                              26-OCT-2000; 2000WO-US029500
                                                                                                                                                                                                          99US-0161532P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
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Best Local Similarity 100.
Matches 14; Conservative
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                                                                                                                                                                                                                                                             Tritz R;
                                                                                                                                                                                                                                                                                     WPI; 2001-300427/31.
                                                                                                                                                                                                                                   (IMMU-) IMMUSOL INC.
                                                                                                                            WO200130362-A2
                                                                                                                                                                                                          26-OCT-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                12-FEB-2004
                                                                                      sapiens
                                                                                                                                                                                                                                                             Robbins JM,
                                                                                                                                                      03-MAY-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 922
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADF37430;
                                                                                                   Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           9
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                                                                                      Homo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADF37430
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cytostatic; antidiabetic; ophthalmological; antiarthritic; antipsoriatic; nephrotropic; gynaecological; angiogenesis-associated condition; cancer; diabetic retinopathy; macular degeneration; neovascular glaucoma; arthritis; psoriasis; endometriosis; angiofibroma; polycystic kidney disease; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The present invention describes a double-stranded short interfering nucleic acid (siNA) that downregulates expression of the vascular endothelial growth factor receptor (VEGRR) gene. Also described: (1) a siNA that downregulates the VEGF gene; (2) kits for in vitro or in vitro delivery of siNA; (3) conjugates and/or complexes of siNA; (4) vectors that express siNA; and (5) single-stranded siNA with similar properties.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           double-stranded interfering nucleic acid, useful e.g. for treatment diagnosis of cancer, downregulates the vascular endothelial growth
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 11-MAR-2002; 2002US-0363124P.
29-MAY-2002; 2002WO-US017674.
06-UUN-2002; 2002US-0386782P.
03-UUL-2002; 2002US-0393348P.
29-JUL-2002; 2002US-0399348P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            29-AUG-2002; 2002US-0406784P.
05-SEP-2002; 2002US-0408378P.
09-SEP-2002; 2002US-0409293P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2002US-0358580P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2002US-00306747
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (RIBO-) RIBOZYME PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 13; Conservative
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Gaps

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Indels

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03-JUL-2002;
29-JUL-2002;
29-AUG-2002;
                                                    20-FEB-2003;
                                      Homo sapiens
                                                         20-FEB-2002;
                                                           11-MAR-2002;
                                                                06-JUN-2002;
                                                                         05-SEP-2002;
                                                                           39-SEP-2002;
                                                                             34-NOV-2002;
                                                                                27-NOV-2002;
      12-FEB-2004
                                               28-AUG-2003
                                    Synthetic
  ADF37677;
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Query Match

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The present invention relates to a method for judging the genotype of hepatitis B viruses (HBV) in which part of the gene sequence of the PreS1 region of HBV is amplified by PCR using labelled primers and the amplified product is hybridized with HBV type A, B, C, D, E, F and G gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Classification of genotype of hepatitis B viruses and primers and probes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Clostridium histolyticum; collagenase; enzymatically active; cleavage; fusion protein; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          specific probes and the label in the PCR product is detected
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0.8%; Score 14; DB 1; Length 19;
100.0%; Pred. No. 9.1e+02;
tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Clostridium histolyticum collagenase PCR primer #1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 19 BP; 9 A; 7 C; 2 G; 1 T; 0 U; 0 Other;
    Mismatches
                                                                                                                                                                                                                                                                                                                                10.
                                                                                                                                                                                                                                                                                                                              SEQ ID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 3; Page 3; 13pp; Japanese.
  0;
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                                                                                                                                                                                                                                                                                                                          Hepatitis B virus PreSl probe,
                                                                                                                                                                                             ADF70750 standard; DNA; 19 BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           14-AUG-2000; 2000JP-00245606
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (GENO-) GENOME SCI KENKYUSHO
                                          180 AGGCATAGACAAGA 193
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CAATCCCAACAAAG 1071
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                                                                                                                                                                                                                                                                                   (first entry)
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  14; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Synthetic.
Clostridium histolyticum.
                                                                                     19 AGGCATAGACAAGA
                                                                                                                                                                                                                                                                                                                                                                     PreS1; HBV; probe; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2003-451644/43
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                Hepatitis B virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                           JP2002355098-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              for the method
                                                                                                                                                                                                                                                                                   12-FEB-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               JP11137256-A
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      14;
                                                                                                                                                                                                                                        ADF70750;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1058
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
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                                                                                                                                                     RESULT 1268
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The present introduction describes a quodice state interface in the present interface acid (siNA) that downregulates expression of the vascular endothelial growth factor receptor (VEGFR) gene. Also described: (1) a candothelial growth factor receptor (VEGFR) gene. Also described: (1) a candother that downregulates the VEGF gene; (2) kits for in vitro or in vivo delivery of siNA; and (5) single-stranded siNA with similar properties. The siNAs have antianglogenic, cytostatic, nephrotropic and gynaecological activities. The siNA are useful for modulating conditions, useful for treating a wide range of angiogenesis-associated conditions, particularly cancers, diabetic retinopathy, macular degeneration, necvascular glaucoma, arthritis, psoriasis, endometriosis, angiofibroma, and polycystic kidney disease. The siNA may also be useful for diagnosis, drug screening, target identification and validation, genetic engineering, studying gene function, and also for gene mapping (e.g. of single-interior or in the condition).
                                                                                                                                       double-stranded short interfering nucleic acid; short interfering nucleic acid; siNA; downregulation; short interfering nucleic acid; siNA; downregulation; candinablat growth factor receptor; VECFR; antiangiogenic; cytostatic; antidiabetic; ophthalmological; antiarthritic; antipsoriatic; nephrotropic; gynaecological; angiogenesis-associated condition; cancer; diabetic retinopathy; macular degeneration; neovascular glaucoma; arthritis; psoriasis; endometriosis; angiofibroma;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       double-stranded interfering nucleic acid, useful e.g. for treatment diagnosis of cancer, downregulates the vascular endothelial growth
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        present invention describes a double-stranded short interfering
                                                                                                 Human VEGFR3 short interfering nucleic acid (siNA) SEQ ID NO:1966.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 14; DB 1; Length 19;
Pred. No. 9.1e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 19 BP; 1 A; 6 C; 3 G; 0 T; 9 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example 3; SEQ ID NO 1966; 207pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mcswiggen J, Beigelman L, Pavco P;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2002US-0386782P-
2002US-0393796P-
2002US-0406784P-
2002US-0408378P-
2002US-0408378P-
2002US-0408378P-
2002US-0408378P-
2002US-0408378P-
2002US-0408378P-
2002US-0408378P-
                                                                                                                                                                                                                                                                          arthritis; psoriasis; endomet:
polycystic kidney disease; ss
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2002WO-US017674.
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                                                        (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     factor receptor gene
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Best Local Similarity
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Gaps

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25-MAY-1999

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The present invention describes an enzymatically active polypeptide (I) derived from a Clostridium histolyticum collagenase with its collagencombining region deleted which specifically recognizes a peptide with the sequence PLGP, and which cleaves the peptide by hydrolysing the peptide bond on C-terminal side of the leucine residue of this sequence and which does not decompose water-insoluble type I collagen. The present sequence represents a PCR primer used in an example from the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Genotyping an alpha-2B, 2A, or 2C adrenergic receptor gene useful for determining whether an individual is at increased risk of developing a disease associated with the corresponding receptor comprises detecting
                                                                                                                                                 New enzymatically active polypeptide and kit containing it - useful for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human; genotyping; alpha-2B; alpha-2A; alpha-2C; adrenergic receptor; polymorphic site; allelic variant; cardiovascular disease; central nervous system disease; adenylyl cyclase; MAP kinase activity; phosphorylation; inositol phosphate; alpha-2BAR; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                 0.8%; Score 14; DB 1; Length 20; 100.0%; Pred. No. 9.5e+02; ative 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                    Sequence 20 BP; 3 A; 3 C; 6 G; 8 T; 0 U; 0 Other;
                                                                                                                                                                                                 Example 1; Page 15; 16pp; Japanese.
                                                                                        (SEGK ) SEIKAGAKU KOGYO CO LTD.
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10-AUG-2000; 2000US-00636259.
19-OCT-2000; 2000US-00692077.
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                                                          97JP-00310887.
                           97JP-00310887
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                                                                                                                                                                      cleaving fusion proteins
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                                                                                                                     WPI; 1999-374377/32.
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Best Local Similarity
Matches 14; Conserv
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(SMAL/) SMALL K M.
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                              12-NOV-1997;
                                                           12-NOV-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The present invention is related to the coding sequence and protein fragments of a human catenin-binding zinc finger protein. The coding
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100.0%; Pred. No. 9.5e+02;
ive 0; Mismatches 0; Indels
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Claim 10; Page 112; 163pp; English.
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Les 14; Conservative
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Homo sapiens
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sequence was isolated from a human kidney cDNA library, but is expressed in most human tissue. The sequences provided by the invention can be used in the diagnosis and treatment of cancer and neurological disorders, and in drug screening to identify compounds capable of the same
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             fragments of a human catenin-binding zinc finger protein. The coding sequence was isolated from a human kidney cDNA library, but is expressed in most human tissue. The sequences provided by the invention can be used in the diagnosis and treatment of cancer and neurological disorders, and in drug screening to identify compounds capable of the same
                                                                                                                                                                                                                                                                                                                                                                                                                                                          Nucleic acid or its fragments, useful for diagnosing and treating cancer and neurological disorders, corresponds to a catenin-binding protein in signal transduction and gene regulatory pathways.
                                                                                             Gaps
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PCR primer; ss.
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                                                                      0.8%; Score 14; DB 1; Length 20; 100.0%; Pred. No. 9.5e+02;
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                                                 Sequence 20 BP; 5 A; 2 C; 8 G; 5 T; 0 U; 0 Other;
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                                                                           100.0%; Pred. ...
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                                                                                                                                                                                                                                                                                                                                                                                                                      Janssens B;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Disclosure; Page 17; 71pp; English.
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                                                                                                           877 GATGACTGTGGGAA 890
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                                                                                                                                                                                AAC88704 standard; DNA; 20
                                                                                                                                2 GATGACTGTGGGAA 15
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                                                                                           Conservative
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Best Local Similarity 100.
Matches 14; Conservative
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                                                                              Local Similarity
es 14; Conserv
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                                                                                                                                                                                                                                                                           screening;
                                                                                                                                                                                                                                                                                              Homo sapiens
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The invention relates to human catenin-binding proteins and their corresponding CDNA molecules which functions in signal transduction and gene regulatory pathways. The invention also provides an isolated and/or recombinant nucleic acid or its functional fragment, homologue or derivative, corresponding to a alpha-catenin binding protein. The invention also relates to a novel human zinc finger protein binding with a member of the a-catulin/vinculin family, preferably with a human isoform of alpha N-catenin (neural form). The invention also relates to the field of drug discovery, diagnosis, prognosis and treatment of cancer and neurological disorders. The present sequence is a primer which is
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Novel recombinant nucleic acids useful for diagnosing, prognosing and/or treating cancer and neurological disorders, corresponds to a protein binding to alpha-catenin protein and with signal transduction function.
                                                                                                                                                                Human; ANC_2H01 protein; catenin-binding protein; signal transduction;
gene regulation; zinc finger protein; alphaN-catenin; drug screening;
therapy; cancer; neurological disorder; cytostatic; neuroprotective;
primer; ss.
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gene regulation; zinc finger protein; alphan-catenin; drug screening;
therapy; cancer; neurological disorder; cytostatic; neuroprotective;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
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                                                                                                                         Human ANC_2H01 cDNA sequencing forward primer, FVR463F.
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Pred. No. 9.5e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Disclosure; Page 66; 160pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0.8%; S
100.0%;
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                                                           (first entry)
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                 Homo sapiens.
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AAD12630;
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ine uigonuciective hybridistes with the incontribute conditions of high stringency and has a sequence comprising 10-40 bp. The internucleoside linkages of the Oligonucleotide comprise at least one phosphorethicate linkage. Hybridisation of the oligonucleotide to the ribonucleic acid inhibits expression of the heparanase, where inhibition of heparanase means at least a 50% reduction in the quality of heparanase in a cell; (1) a method of inhibiting expression of a heparanase in a cell; (2) a composition comprising the above oligonucleotide in an amount effective to inhibit the expression of the heparanase in the cell and a carrier; and (3) a method of tracating a cumour in a subject comprises administering to the subject an amount of the above oligonucleotide effective to inhibit expression of a heparanase in the subject comprises administering to the subject an amount of the above oligonucleotide effective to inhibit expression of a heparanase control of the colligonucleotide enterapy, and can be used for preparing a cutivity, can be used in gene therapy, and can be used for preparing a composition for treating tumours. The present sequence represents a human control of the paranase phosphorothicate antisense oligonucleotide, which is used in
                                                                                                                                                                                                                                                                                          The present invention describes an oligonucleotide having a sequence complementary to a sequence of ribonucleic acid encoding a heparanase. The oligonucleotide hybridises with the ribonucleic acid under conditions
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /mod_base= OTHER
/mote= "This oligonucleotide has a phosphorothioate
/note= "This oligonucleotide (2-MoB) wings at the 5'
backbone and 2-0-methoxyethyl (2-MoB) wings at the 5'
and 3' ends, which are 5 nucleotides in length. Also all
cytidine residues are 5-methylcytidines"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Vascular endothelial growth factor receptor 1, VEGF receptor; VEGFR, inhibitor; cytostatic; antirheumatic; antiarthritic; antiangiogenic; antiarthritic; antiangiogenic; antiarisense gene therapy; hyperproliferative disorder; cancer; rheumatoid arthritis; angiogenesis; infection; inflammation; tumour formation; phosphorothioate; 2'-0-methoxyethyl; 2'-MOE; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mouse VEGFR-1 chimeric phosphorothioate oligonucleotide SEQ ID NO:143
                                                                                                                                        New oligonuclectide having a sequence complementary to a sequence of ribonucleic acid encoding a heparanase, useful for preparing a composition for treating tumor.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 14; DB 1; Length 20;
Pred. No. 9.5e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 20 BP; 6 A; 8 C; 6 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          the exemplification of the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           100.0%; Pred. ...
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                                                                                                                                                                                                                                                   Claim 7; Page 32; 48pp; English.
(UYCO ) UNIV COLUMBIA NEW YORK.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         BP,
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Best Local Similarity 100.
Matches 14; Conservative
                                                                                                WPI; 2003-201558/19
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Synthetic.
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                                                    Stein C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             RESULT 1276
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The invention relates to human catenin-binding proteins and their corresponding cDNA molecules which functions in signal transduction and gene regulatory pathways. The invention also provides an isolated and/or recombinant nucleic acid or its functional fragment, homologue or derivative, corresponding to a alpha-catenin binding protein. The invention also relates to a novel human zinc finger protein binding with a member of the a-catulin/vinculin family, preferably with a human isoform of alpha N-catenin (neural form). The invention also relates to the field of drug discovery, diagnosis, prognosis and treatment of cancer and neurological disorders. The present sequence is a primer which is
                                                                                                                                                                                                                                                                                                                                                         Novel recombinant nucleic acids useful for diagnosing, prognosing and/or treating cancer and neurological disorders, corresponds to a protein binding to alpha-catenin protein and with signal transduction function.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human; heparanase; phosphorothioate; antisense oligonucleotide; cytostatic; gene therapy; tumour; ss.
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0.8%; Score 14; DB 1; Length 20;
Best Local Similarity 100.0%; Pred. No. 9.5e+02;
Matches 14; Conservative 0; Mismatches 0; Indels
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                                                                                                                                                                                                            (VLAA-) VLAAMS INTERUNIVERSITAIR INST BIOTECHNOG
                                                                                                                                                                                                                                                                Janssens B;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          used for sequencing human ANC_2H01 cDNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Disclosure; Page 66; 160pp; English.
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/*tag= a
/mod_base= OTHER
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                                                                                                                18-MAY-2000; 2000WO-EP004535.
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modified_base
             WO200147954-A2
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                                                                                                                                                                  23-DEC-1999;
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                                                                 05-JUL-2001
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RESULT 1275

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ABZ22802

Homo

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Gaps .. 0

0; Indels

12-SEP-2002; 2002WO-US029148, 13-SEP-2001; 2001US-00953318. (ISIS-) ISIS PHARM INC Bennett CF, Watt AT; WPI; 2003-301004/29. 20-MAR-2003

New antisense oligonucleotide targeted to a nucleic acid encoding vascular endothelial growth factor receptor-1, useful for diagnosing or treating cancer, rheumatoid arthritis, or diseases or conditions involving angiogenesis.

Claim 3; Page 86; 150pp; English.

transplant and severable a compound (1) or or uncleobases in length targeted to a nucleic acid molecule encoding vascular endothelial growth factor receptor-1 (VEGRP-1), where the compound inhibits the expression of VEGRP-1 or with an 8-nucleobase portion of an active site on the nucleic acid encoding C VEGRP-1 or with an 8-nucleobase portion of an active site on the nucleic acid molecule encoding VEGRP-1. Also described: (1) a composition or comprising (C) and a carrier or diluent; (2) inhibiting the expression of VEGRP-1 in cells or tissues by contacting the cells or tissues with (C) so that the expression of VEGRP-1 is inhibited; and (3) treating an animal having a disease or condition associated with VEGRP-1 is unlial having a disease or condition associated with VEGRP-1 is inhibited. (C) has antiangiogenic, antirheumatic, antiarthritic, cytostatic and antiinflammatory activities, and can be used in antisense gene therapy. The antisense compounds are useful for modulating the expression of VEGRP-1, such as hyperproliferative disorders of with the expression of VEGRP-1, such as hyperproliferative disorders consideration or theraptures, prophylaxis, e.g. to prevent or callay infection, therapeutics, prophylaxis, e.g. to prevent or delay infection, confitions involving the pathwart formation, as research reagents and kits, and in the pharmarian antisense compounds when the encounter therapeut presents sequence represents a mouse VEGRP-2 chimarian antisense or hyperparties members of a biological membrane antisense confit antisense consequence antisense consequence antisense confit and in a several confit and in a particular antisense of a biological antisense confit and antisense present invention describes a compound (C) 8-50 nucleobases in length shosphorothioate antisense oligonucleotide, which is used in an example from the present invention

Sequence 20 BP; 7 A; 3 C; 5 G; 5 T; 0 U; 0 Other;

.; 0 0.8%; Score 14; DB 1; Length 20; 100.0%; Pred. No. 9.5e+02; Live 0; Mismatches 0; Indels Query Match 0.8 Best Local Similarity 100. Matches 14; Conservative

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Gaps

539 CCATCTTTGACAAG 552 ccarcificacaag s 18

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g

RESULT 1277

ABZ93277 standard; DNA; 20 BP ABZ93277;

Human oligonucleotide sequence

17-OCT-2003 (first entry)

Human, antisense, lung dysfunction, nasal airway dysfunction, antiniflammatory steroid; ubiquinone; antiniflammatory; antiallergic; antiasthmatic; hypotensive; cymostatic; gene therapy, antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy; lung inflammation; respiratory disease; ds.

Homo sapiens

The invention relates to a novel pharmaceutical composition, which has a first active agent comprising an oligonucleotide antisense to the initiation coodon, coding regalon, 5' or 3' end genomic flanking regions, 5' and 3' intron-exon junctions, or regions within 2-10 nucleotides of junctions of genes encoding a polypeptide associated with lung and/or nasal airway dysfunction and a second active agent comprising an entilnflammatory steroid and ubiquinone. A composition of the invention has antiinflammatory, antiallargic, antiasthmatic, hypotensive, immunosuppressive, and cytostatic activity. The composition may have a tree in antisense gene therappy. The composition is useful for treating or preventing a respiratory, lung or malignant disease or condition, also for enhancing the prophylactic or therapeutic respiratory effect of an antiinflammatory steroid in a subject, for reducing or depleting levels of, or reducing sensitivity to adenosine, reducing levels of adenosine receptor, producing bronchodilation, increasing levels of ubiquinone or lung surfactant in a subject, for reading levels of ubiquinone or lung surfactant in a subject, stissue, or treating bronchoconstriction, when it is a subject is the subject in the subject in the subject in the subject is the subject in t Pharmaceutical composition for treating ailments associated with impaired lung inflammation, lung allergies, or a respiratory disease or condition. Note: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or Gaps Katz E, Pabalan J, Aguilar D; .; 0 0.8%; Score 14; DB 1; Length 20; 100.0%; Pred. No. 9.5e+02; vative 0; Mismatches 0; Indels Sequence 20 BP; 6 A; 8 C; 3 G; 3 T; 0 U; 0 Other; ftp.wipo.int/pub/published_pct_sequences Disclosure; SEQ ID NO 8519; 872pp; English. Li Y, Sandrasagra A, Ka Tang L, Shahabuddin S; 23-APR-2002; 2002WO-US013135. 24-APR-2001; 2001US-0286137P. (EPIG-) EPIGENESIS PHARM INC. 1087 GIGGIGACACIGIG 1100 Conservative WPI; 2003-229219/22. Similarity WO200285308-A2. Mac Local Sim. 14; 31-OCT-2002 ubiquinone. Miller S, Nyce JW. Query Match Best Loca Matches g

14 Greereachere 1

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ABD29507 standard; DNA; 20 ABD29507; ABD29507/

RESULT 1278

BP.

AA664176-derived oligonucleotide SEQ ID 8519.

(first entry)

29-JUL-2004

Human, antisense, bronchoconstriction; allergy, hyposecretion; pain; respiratory tract inflammation, adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antiinflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory distress syndrome; allergic rhinitis; pulmonary hypotremaion; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis; pulmonary transplantation rejection; ss; primer. 22-APR-2004 (first entry)

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This invention describes a novel composition (a) a first active agent, comprising oligonucleotides, effective for alleviating comprising oligonucleotides, effective for alleviating concentration, respiratory tract inflammation, allergies and reducing adenosine sensitivity, levels of adenosine (A) or (A) receptors, contractant depletion or hyposecretion, when administered to a mammal. The corporation of a target polypeptide associated with lung airway or lung expression of a target polypeptide associated with lung airway or lung dysfunction or cancer and can be anti-sense to the corresponding mRNA. The invention also describes a kit, that comprises: (a) a delivery device, in separate containers (b) the oligonucleotides, (c) instructions for adding a carrier and for use of the kit. The composition of the invention has antiallergic, antialflammatory, antiathmanetory, antiathmanetory, antiathmanetory, antiathmanetory, is a composition comprises oligo and is administered to reduce the production or availability, or to increase the degradation of the target mRNA or to reduce the amount of target polypeptide present in the lungs. The pulmonary obstruction, and/or bronchoconstriction and/or lung inflammation, allergies and/or bring the proproduction are associated inflammation, allergies and/or surfactant hypoproduction are associated inflammation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            inflammation, allergies, asthma, impeded respiration, respiratory distress syndrome, pain, cystic fibrosis, allergic rhinitis, pulmonary hypertension, emphysema, chronic obstructive pulmonary disease, pulmonary
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The reduced adenosine content of the anti-sense oligos corresponding to thymidines present in the target RNA serves to prevent the breakdown of the oligonucleotides into products that free adenosine into the system
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          e.g., lung, brain, heart, kidney, etc, tissue environment and thereby, prevent any unwanted effects due to it
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           hypertension, emphysema, chronic obstructive pulmonary disease, pulmor
transplantation rejection, pulmonary infections, bronchitis or cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                           oligonucleotide containing less percentage of adenosine, targeted to nucleic acids associated with lung airway or lung dysfunction, and
                                                                                                                                                                                                                                                                                                                                                                                                                       Pharmaceutical composition for treating asthma, has antisense
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 20 BP; 6 A; 8 C; 3 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 15; SEQ ID NO 8519; 763pp; English.
                                                                                                                                                                                                          24-APR-2001; 2001US-0286036P.
                                                                                                                                                                23-APR-2002; 2002WO-US013143
                                                                                                                                                                                                                                                           (EPIG-) EPIGENESIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 bronchodilating agent
                                                                                                                                                                                                                                                                                                                                                                             WPI; 2003-093058/08.
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Tang L,
                                                                  WO200285309-A2
                      Homo sapiens.
                                                                                                                 31-OCT-2002
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Miller S,
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                          Gaps
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0
Score 14; DB 1; Length 20;
Pred. No. 9.5e+02;
0; Mismatches 0; Indels
      100.001
100.001
100.001
    0.88;
                            Conservative
                 Similarity
    Query Match
Best Local Simi
Matches 14;
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1087 GTGGTGACACTGTG 1100
                         Greergacacrere 1
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ADJ34005 standard; DNA; 20 BP. ADJ34005; RESULT 1279 ADJ34005, H XXXX

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cytostatic; antiinflammatory; antimicrobial; antisense gene therapy; kinase inhibitor; hyperproliferative disorder; cancer; non-small cell lung cancer; oesophageal cancer; infection; inflammation; tumour; human; phosphorothioate; 2'-0-methoxyethyl; ss.
                    polo-like kinase; polo-like kinase inhibitor; antisense oligonucleotide;
        Human polo-like kinase antisense oligonucleotide SEQ ID NO:65.
                                                                                                /note= "phosphorothioate linkages"
                                                                                                                   /note= "2'-0-methoxyethyls"
16. .20
                                                                          Location/Qualifiers
                                                                                            base= OTHER
                                                                                                                   OTHER
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                                                                                             /mod
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modified_base
                                                                                                       modified_base
                                                         sapiens.
                                                                Synthetic.
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Aguilar D;

Pabalan J,

Katz E,

Sandrasagra A, Ka ,, Shahabuddin S;

WO2004011610-A2 05-FEB-2004

/mod_base= OTHER
/note= "2'-0-methoxyethyls"

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/*tag=

modified base

25-JUL-2003; 2003WO-US023413.

30-JUL-2002; 2002US-00209405.

(ISIS-) ISIS PHARM INC

Freier SM; WPI; 2004-143840/14. Wyatt JR,

New antisense compounds targeted to nucleic acid molecules encoding pololike kinase, useful for treating diseases associated with aberrant expression of polo-like kinase, e.g. non-small cell lung cancer or esophageal cancer.

Example 15; SEQ ID NO 65; 138pp; English.

The present invention describes a compound (I) of 8-80 nucleobases in length targeted to a nucleic acid molecule encoding polo-like kinase, where (I) specifically hybridises with nucleic acid molecule encoding copol-like kinase and inhibits the expression of polo-like kinase, or specifically hybridises with at least an 8-nucleobase portion of a specifically hybridises with at least an 8-nucleobase portion of preferred target region on a nucleic acid molecule encoding polo-like kinase. Also described: (I) a composition comprising (I) and a composition comprising (I) and a composition comprising (I) and a composition contacting the expression of polocitismes in cells or tissues comprising administering to the animal associated with polo-like kinase comprising administering to the animal contacting associated with polo-like kinase comprising a nucleic and (1) so that expression of polo-like kinase is inhibitated; and (4) screening for an antisense compound comprising at least an 8-nucleobase portion which is compoundes comprising at least an 8-nucleobase portion which is compounded at least encoding polo-like kinase with one or more candidate antisense compounds which inhibits the expression of a nucleic acid molecule encoding polo-like kinase (I) has cytostatic, antiful amanery, and antisense (I) has cytostatic, antiful amanery, and antisense (I) has cytostatic, antiful amanery and antisense (I) antifit and antisense compounds of a nucleic acid molecule encoding polo-like kinase (I) has cytostatic, antiful amanery and antisense compounds and antisense compounds and antisense compounds of a nucleic acid molecule encoding polo-like kinase (I) has cytostatic, antiful amanery and antisense compounds antisense compounds antisense compounds antisense compounds antisense compounds ant antientlammatory and antimicrobial activities, and can be used in antiennes gene therapy, and as a kinase inhibitor. The antienne oligonucleotides or compounds (I) can be used for inhibiting the expression of polo-like kinase, and for treating diseases or conditions associated with aberrant expression of polo-like kinase, e.g. hyperproliferative disorder such as cancer, including non-small cell lung

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·,
                                                  human
                                                                                                                                                                                                                                                                                                                                                                                                                          cytostatic; antidiabetic; immunomodulator; cardiant; neuroprotective; gene therapy; endothelial specific molecule-1; ESM-1; ESM-1; ESM-1 related disorder; diabetes; cancer; ischaemia; reperfusion injury; angiogenic disorder; immunological disorder; cardiovascular disorder; neurological disorder; se.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New antisense compound, having a sequence targeted to a nucleic acid encoding endothelial specific molecule-1 (ESM-1), useful for preparing a composition for treating e.g., diabetes, cancer or cardiovascular
cancer or oesophageal cancer. The antisense compounds are also useful research reagents and kits, or in diagnostic, therapeutic and prophylactic applications, e.g. to prevent or delay infection, inflammation or tumour formation. The present sequence represents a hupolo-like kinase chimeric phosphorothioate antisense oligonucleotide, which is used in an example from the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /note= "OTHER= phosphorothioate backbone. All cytidine residues are 5-methylcytidines"
                                                                                                                                                                        Gaps
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/note= "OTHER= 2'-methoxyethyl (2'-MOE) nucleotides"
16. .20
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                                                                                                                                     Query Match 0.8%; Score 14; DB 1; Length 20; Best Local Similarity 100.0%; Pred. No. 9.5e+02; Matches 14; Conservative 0; Mismatches 0; Indels
                                                                                                         Sequence 20 BP; 5 A; 4 C; 4 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                             Human ESM-1 antisense oligonucleotide segid 544.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 3; SEQ ID NO 544; 555pp; English
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/mod_base= OTHER
                                                                                                                                                                                                                                                                                                       ADL58295 standard; DNA; 20 BP.
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                                                                                                                                                                                                 132 GATGAAGAAGATCA 145
                                                                                                                                                                                                                                                                                                                                                                 03-JUN-2004 (first entry)
                                                                                                                                                                                                                             GATGAAGAAGATCA 2
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The invention describes a new antisense compound, having a sequence

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comprising 8-30 bp targeted to a nucleic acid encoding endothelial specific molecule-1 (ESM-1), that specifically hybridises with the nucleic acid ESM-1 and inhibits its expression. Also described are: a composition; inhibiting the expression of ESM-1 in cells or tissues; and treating an animal having a disease or condition associated with ESM-1. The compound is useful for preparing a composition for treating diabetes, cancer, ischaemia or reperfusion injury, or angiogenic, immunological, cardiovascular or neurological disorder. This sequence represents an antisense oligonuclectide that can be used to modulate expression of endothelial specific molecule-1 (ESM-1).
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/note= "OTHER= phosphorothioate backbone. All cytidine
residues are 5-methylcytidines"
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100.0%; Pred. No. 9.5e+02;
ive 0; Mismatches 0; Indels
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/note= "OTHER= 2'-methoxyethyl (2'-MOE)
16. .20
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1es 14; Conservative
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Location/Qualifiers
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Matches 14; Conservative
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ADL58665/c
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                                        comprising 8-30 bp targeted to a nucleic acid encoding endothelial specific molecule-1 (ESM-1), that specifically hybridises with the nucleic acid ESM-1 and inhibits its expression. Also described are: a composition, inhibiting the expression of ESM-1 in cells or tissues; and treating an animal having a disease or condition associated with ESM-1. The compound is useful for preparing a composition for treating diabetes, cancer, ischaemia or reperfusion injury, or anglogenic, immunological, cardiovascular or neurological disorder. This sequence represents an antisense oligonoclecide that can be used to modulate expression of
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/note= "OTHER= phosphorothioate backbone. All cytidine
residues are 5-methylcytidines"
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                                  The invention describes a new antisense compound, having a sequence
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100.0%; Pred. No. 9.5e+02;
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               Claim 3; SEQ ID NO 1354; 555pp; English
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Best Local Similarity
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/note= "OTHER= phosphorothioate backbone. All cytidine
residues are 5-methylcytidines"
New antisense compound, having a sequence targeted to a nucleic acid encoding endothelial specific molecule-1 (ESM-1), useful for preparing composition for treating e.g., diabetes, cancer or cardiovascular
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100.0%; Pred. No. 9.5e+02;
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/mod_base= OTHER
/note= "OTHER= 2'-methoxyethyl (2'-MOE)
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Griggs DW

Weinstein EJ,

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The invention describes a new antisense compound, having a sequence comparising 8-30 bp targeted to a nucleic acid encoding endothelial specific molecule-1 (ESM-1), that specifically tybridises with the nucleic acid ESM-1 and inhibits its expression. Also described are: a composition; inhibiting the expression of ESM-1 in cells or tissues; and treating an animal having a disease or condition associated with ESM-1. The compound is useful for preparing a composition for treating diabetes, cancer, ischaemia or reperfusion injury, or angiogenic, immunological, cardiovascular or neurological diaborder. This sequence represents an antisanse oligonucleotide that can be used to modulate expression of endothelial specific molecule-1 (ESM-1).
                                                              New antisense compound, having a sequence targeted to a nucleic acid encoding endothelial specific molecule-1 (ESM-1), useful for preparing a composition for treating e.g., diabetes, cancer or cardiovascular
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                                                                                                                                           Claim 3; SEQ ID NO 914; 555pp; English.
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19-AUG-2003; 2003WO-US025833

WO2004021978-A2

18-MAR-2004

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Score 14; DB 1; Length 20; Pred. No. 9.5e+02; 0; Mismatches 0; Indels

100.0%; Pred. ...

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The invention describes a new antisense compound, having a sequence comprising 8-30 bp targeted to a nucleic acid encoding endothelial specific molecule. (ESM-1), that specifically hybridises with the nucleic acid ESM-1 and inhibits its expression. Also described are: a composition; inhibiting the expression of ESM-1 in cells or tissues; and treating an animal having a disease or condition associated with ESM-1. The compound is useful for preparing a composition for treating diabetes, cancer, ischaemia or reperfusion injury, or angiogenic, immunological, cardiovascular or neurological disorder. This sequence represents an antisense oligoniclectide that can be used to modulate expression of
                                                                                                                                        New antisense compound, having a sequence targeted to a nucleic acid encoding endothelial specific molecule-1 (ESM-1), useful for preparing a composition for treating e.g., diabetes, cancer or cardiovascular
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16. .20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 20 BP; 9 A; 6 C; 1 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human ESM-1 antisense oligonucleotide seqid 1095.
                                                                                                                                                                                                                                                                                                                                                                                                                                            endothelial specific molecule-1 (ESM-1)
                                                                                                                                                                                                                           Claim 3; SEQ ID NO 673; 555pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADL58846 standard; DNA; 20 BP.
19-AUG-2002; 2002US-0404495P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match 0.8
Best Local Similarity 100.
Matches 14; Conservative
                                    (PHAA ) PHARMACIA CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           *tag=
                                                                     Griggs
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                                                                      Weinstein EJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Key
modified_base
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                                                                                                                                                                                            disorder
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RESULT 1285
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/note= "OTHER= phosphorothioate backbone. All cytidine
residues are 5-methylcytidines"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /note= "OTHER= 2'-methoxyethyl (2'-MOE) nucleotides"
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/note= "OTHER= 2'-methoxyethyl (2'-MOE) nucleotides"
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Griggs DW;
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                                                                                                     (PHAA ) PHARMACIA CORP
                                                                                                                                             WPI; 2004-248358/23.
                                                                                                                                                                                                                                                                                                                                                                                                     Local Similarity
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                 WO2004021978-A2
                                                                                                                         Weinstein EJ,
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                                                                                                                                                                                                                                                                                                                                                                                                                14;
                                        18-MAR-2004.
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                                                                                                                                                                                                                                                                                                                                                                                                                                     593
                                                                                                                                                                                                                                                                                                                                                                                                                                                          13
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                                                                                                                                                                                                The invention describes a new antisense compound, having a sequence comprising 8-30 bp targeted to a nucleic acid encoding endothelial specific molecule-1 (ESM-1), that specifically Mybridises with the nucleic acid ESM-1 and inhibits its expression. Also described are: a composition; inhibiting the expression of ESM-1 in cells or tissues; and treating an animal having a disease or condition associated with ESM-1. The compound is useful for preparing a composition for treating diabetes, cancer, ischaemia or reperfusion injury, or angiogenic, immunological, cardiovascular or neurological disorder. This sequence represents an antisense oligonomicleotied that can be used to modulate expression of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      cytostatic; antidiabetic; immunomodulator; cardiant; neuroprotective; gene therapy; endothelial specific molecule-1; ESM-1; ESM-1shated disorder; diabetes; cancer; ischaemia; reperfusion injury; angiogenic disorder; immunological disorder; cardiovascular disorder; neurological disorder; se.
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                                                                                                                          New antisense compound, having a sequence targeted to a nucleic acid encoding endothelial specific molecule-1 (ESM-1), useful for preparing composition for treating e.g., diabetes, cancer or cardiovascular
                                                                                                                                                                                                                                                                                                                                                                          Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /mod_base= OTHER
/note= "OTHER= 2'-methoxyethyl (2'-MOE) nucleotides"
                                                                                                                                                                                                                                                                                                                                                                         0;
                                                                                                                                                                                                                                                                                                                                                     Length 20;
                                                                                                                                                                                                                                                                                                                                                                         0; Indels
                                                                                                                                                                                                                                                                                                                                 Sequence 20 BP; 10 A; 5 C; 1 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                     DB 1; Le
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human ESM-1 antisense oligonucleotide segid 1096.
                                                                                                                                                                                                                                                                                                                                                                          Migmatches
                                                                                                                                                                                                                                                                                                               endothelial specific molecule-1 (ESM-1).
                                                                                                                                                                                                                                                                                                                                                      Score 14;
Pred. No.
                                                                                                                                                                                Claim 3; SEQ ID NO 1095; 555pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                             100.0%; Pre
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                       19-AUG-2003; 2003WO-US025833
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                                             19-AUG-2002; 2002US-0404495P
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                                                                                      Griggs DW;
                                                                                                                                                                                                                                                                                                                                                                           Conservative
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                                                                (PHAA ) PHARMACIA CORP.
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                                                                                                          WPI; 2004-248358/23
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Les 14; Conserv
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modified base
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                                                                                      Weinstein EJ,
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   18-MAR-2004
                                                                                                                                                                                                                                                                                                                                                                                                593
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                                                                                                                                                                                                                                                                                                                                                       Query Match
                                                                                                                                                               disorder.
                                                                                                                                                                                                                                                                                                                                                                                                                                                    RESULT 1286
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ss; antidiabetic; immunosuppressive; cardiovascular; antilipemic; antiarteriosclerotic; hepatotropic; litholytic; anorectic; enceptective; vasotropic; antisense; gene therapy; Farnesoid X receptor; diabetes; immunological disorder; cardiovascular disorder; dyslipidemia; atherosclerosis; high density lipoproteni, low density lipoproteni, hypercholesterolemia; gallstones; hypertriglyceridemia; obesity; neurological disorder; ischemia; reperfusion; diagnostics; prophylaxis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New antiBense compound, having a sequence targeted to a nucleic acid encoding endothelial specific molecule-1 (ESM-1), useful for preparing a composition for treating e.g., diabetes, cancer or cardiovascular
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
/note= "OTHER= 2'-methoxyethyl (2'-MOE) nucleotides'
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Pred. No. 9.5e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         claim 3; SEQ ID NO 1096; 555pp; English.
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100.0%; Pre
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                                                                                                                                                                                                                                                                                                                        19-AUG-2002; 2002US-0404495P
                                                                                                                                                                                                                                            19-AUG-2003; 2003WO-US025833
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ADO53870 standard; DNA; 20 BP
25-SEP-2002; 2002US-0413588P
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                                                                                                                                                                                                                                                                                                                                           25-SEP-2003; 2003WO-US030353
                                                                                                                                                                                              6 ACACCCTCACAGG 19
                                                                                                                                                                                                                                            15-JUL-2004 (first entry)
                                                                                                                                                                           14; Conservative
         (PHAA ) PHARMACIA CORP
                                                                                                                                                                                                                                                                                                                                                                (PHAA ) PHARMACIA CORP
                                                       gallstones or obesity.
                             WPI; 2004-347928/32
                                                                                                                                                                     Best Local Similarity
                                                                                                                                                                                                                                                                                                                        WO2004030750-A1
                                                                                                                                                                                                                                                                                                              Homo sapiens
                                                                                                                                                                                                                                                                                                                                  15-APR-2004
                                                                                                                                            invention
                                                                                                                                                                 Query Match
                                                                                                                                                                                                                                  ADO53870
                   Kane CD;
                                                                                                                                                                                                                                                                                                                                                                           Kane CD;
                                                                                                                                                                                                              RESULT 1288
                                                                                                                                                                           Matches
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The invention relates to an antisense compound 8-30 nucleobases in length targeted to a nucleic acid molecule encoding Farnesoid X receptor (FXR), where the antisense compound specifically hybridizes with and inhibits the expression of FXR. The composition and methods are useful for inhibiting the expression of FXR (Farnesoid X receptor) in cells or tissues, or for treating diseases or conditions associated with FXR, such as diabetes, immunological disorders, cardiovascular disorders, e.g. dyslipidemia and its symptoms, atherosclerosis, low HDL (high density lipoprotein), elevated LDL (low density lipoprotein) or hypercholesterolemia, gallstones, hyperthighyeridemia, obesity, hypercholesterolemia, gallstones, hyperthighyeridemia, obesity, neurological disorders, or ischemia/reperfusion injury. In addition, the composition is used for diagnostics, prophylaxis, or as research reagents or kits. This sequence corresponds to an antisense oligonucleotide of the
                                                                                              Farnesoid X Receptor (FXR) or for treating diseases associated with FXR, e.g. diabetes, immunological disorders, cardiovascular disorders,
                                                                    New antisense oligonucleotides useful for modulating expression of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human, Ephrin-B2; ss; antisense oligonucleotide;
phosphorothioate linkage; 2'-O-methoxyethyl sugar moiety;
5-methylcytosine; hyperproliferative disorder; cancer; cytostatic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0.8%; Score 14; DB 1; Length 20; 100.0%; Pred. No. 9.5e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human Ephrin-B2 DNA antisense oligonucleotide #14.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 20 BP; 5 A; 9 C; 2 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              100.0%; Prec. ...
                                                                                                                                                                                            Claim 4; SEQ ID NO 1243; 150pp; English
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les 14; Conservative
                                                                                                                       e.g. diabetes, immunol gallstones or obesity.
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                      2004-347928/32
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      26-AUG-2004
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RESULT 1289
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                                                                                                                                                                                                                                                                                                                                                                         targeted to a nucleic acid molecule encoding Parnesoid X receptor (FXR), where the antisense compound specifically hybridizes with and inhibits the expression of FXR. The composition and methods are useful for inhibiturg the expression of FXR. (Farnesoid X receptor) in cells or tissues, or for treating diseases or conditions associated with FXR, such as diabetes, immunological disorders, cardiovascular disorders, e.g. dyslipidemia and its symptoms, atherosclerosis, low HDL (high density lipoprotein), elevated LDL (low density lipoprotein) or hypercholesterolemia, gallstones, hypertriglyceridemia, obesity, composition is used for diagnostics, prophylaxis, or as research reagents or fits. This sequence corresponds to an antisense oligonucleotide of the
                                                                                                                                                                                                                                                                                                                                                          invention relates to an antisense compound 8-30 nucleobases in length
                                                                                                                                                                               New antisense oligonucleotides useful for modulating expression of Farnesoid X Receptor (FXR) or for treating diseases associated with FXR, e.g. diabetes, immunological disorders, cardiovascular disorders,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ss; antidiabetic; immunosuppressive; cardiovascular; antilipemic; antiatteriosclerotic; hepatotropic, litholytic; anorectic; neuroprotective; vasotropic; antisense; gene therapy; Farnesoid X receptor; diabetes; immunological disorder; cardiovascular disorder; dyslipidemia; atherosclerosis; high density lipoprotein; low density lipoprotein; hypertriglyceridemia; obesity; neurological disorder; gallstones; hypertriglyceridemia; obesity; neurological disorder; ischemia; reperfusion; diagnostics; prophylaxis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Farnesoid X receptor gene expression antisense inhibitory oligo #1243.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ·.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 14; DB 1; Length 20;
Pred. No. 9.5e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 20 BP; 4 A; 9 C; 2 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         100.08; Preu. ...
                                                                                                                                                                                                                                                                                                  Claim 4; SEQ ID NO 1175; 150pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0.88;
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The invention relates to a compound targeted to a nucleic acid molecule

Example 15; SEQ ID NO 26; 69pp; English.

disorder, e.g. cancer.

WPI; 2004-440339/41.

New oligonucleotide compound that inhibits expression of Ephrin-B2, useful for preparing a composition for treating hyperproliferative

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Gaps

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0; Indels

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AAX09121-X10268 are allele-specific oligonucleotide primers used in the isolation of various biallelic polymorphic markers found in the human genome (represented in AAX10269-X1937). These primers can be used in a method for determining polymorphic forms in an individual for use in e.g. forensics, paternity testing or for phenotypic typing for diseases such as agammaglobulinemia, diabetes insipidua. Lesch-Nyhan syndrome, muscular dystrophy, Wiskott-Aldrich syndrome, Fabry's disease, familial hypercholesterolemia, polycystic kidney disease, hereditary hapercholesterolemia, polycystic kidney disease, hereditary haemorrhagic telangiectasia, familial colonic polyposis, Ehlers-Danlos syndrome, osteogenesis imperfecta, acute intermittent porphyria,
                                                                                                                                                                                                                                                                                                                                ö
encoding the human Ephrin-B2 polypeptide. The compound is an antisense oligonucleotide that specifically hybridises with the nucleic acid and inhibite expression of the polypeptide. The antisense oligonucleotide comprises at least one modified internucleoside linkage i.e. a phosphorothioate linkage, at least one modified sugar moiety, preferably a 2-0-methoxyethyl sugar moiety, or at least one modified nucleobase comprising a 5-methylcytosine. The antisense compounds are useful for modulating the expression of the human Ephrin-B2 polypeptide and in preparation of a composition for treating hyperroliferative disorders, e.g. cancer. This sequence represents an antisense oligonucleotide targeted to DNA encoding the human Ephrin-B2 polypeptide of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Polymorphism; biallelic; human; forensic; paternity testing; disease; detection; phenotypic typing; characteristic; infection; hereditary; autoimmune disease; cancer; inflammation; drug; therapy; medicament;
                                                                                                                                                                                                                                                                                                                                Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New isolated nucleic acid segments from the human genome - used for determining polymorphic forms for use in e.g. forensics, paternity testing or phenotypic typing for disease.
                                                                                                                                                                                                                                                                                                                                  .,
                                                                                                                                                                                                                                                                                             DB 1; Length 20; 9.5e+02;
                                                                                                                                                                                                                                                                                                                                0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human biallelic polymorphic marker upstream primer #42.
                                                                                                                                                                                                                                                     Sequence 20 BP; 5 A; 4 C; 4 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                            100.0%; Pred. worthive 0; Mismatches
                                                                                                                                                                                                                                                                                              Score 14;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    RES.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 15; Page 51; 310pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (WHED ) WHITEHEAD INST BIOMEDICAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Hudson T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          21 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        treatment; marker; primer; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            97WO-US020313.
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                                                                                                                                                                                                                                                                                                0.8%;
                                                                                                                                                                                                                                                                                                                                                                          860
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first entry)
                                                                                                                                                                                                                                                                                                                  Best Local Similarity 100.
Matches 14; Conservative
                                                                                                                                                                                                                                                                                                                                                                        847 TACCTGGACAAGGA
                                                                                                                                                                                                                                                                                                                                                                                                             14 TACCTGGACAAGGA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAX09162 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Wang D,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 1998-286974/25.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            05-NOV-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                06-NOV-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WO9820165-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      24-MAR-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       14-MAY-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Lander ES,
                                                                                                                                                                                                                         invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAX09162;
                                                                                                                                                                                                                                                                                                  Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       RESULT 1290
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAX09162
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autoimmune diseases, inflammation, cancer, diseases of the nervous system, infection by pathogenic microorganisms, and characteristics such as longevity, appearance (e.g. baldness, obesity), strength, speed, endurance, fertility, and susceptibility or receptivity to particular drugs on therapeutic treatments. The isolated polymorphic nucleic acid segments can also be used to produce medicaments for the treatment or prophylaxis of such diseases
                                                                                                                                                                                                                                                                                                                                                                                                                                                           ATP binding cassette; ABC transporter; ABCR; Stargardt Disease; therapy; Fundus Flavimaculatus; age-related macular degeneration; diagnosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ΡM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Retina-specific ATP-binding cassette transporter and DNA - useful for, e.g. diagnosis and treatment of macular degeneration, such as in Stargardt Disease, Fundus Flavimaculatus and age-related degeneration.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Li Y;
Smallwood
                                                                                                                                                                                           Gaps
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Singh N,
                                                                                                                                                            Score 14; DB 1; Length 21;
Pred. No. 9.8e+02;
                                                                                                                                                                                             Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 21 BP; 8 A; 3 C; 6 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                               PCR primer ABCR.EXON7:F for ABCR coding sequence.
                                                                                                                                                                                             .;
0
                                                                                                                               Sequence 21 BP; 7 A; 4 C; 6 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Leppart M,
Shroyer NF,
                                                                                                                                                                                             Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (BAYU ) BAYLOR COLLEGE MEDICINE. (UVJO ) UNIV JOHNS HOPKINS. (USSH ) US DEPT HEALTH & HUMAN SERVICES. (UTAH ) UNIV UTAH.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Dean M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Anderson KL, Dean M,
athans J, Rattner A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 41; Page 27; 79pp; English.
                                                                                                                                                                         100.0%; Pre
                                                                                                                                                                                                                                                                                                                                      BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   98WO-US003895.
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                                                                                                                                                                 0.8%;
                                                                                                                                                                                                                           714 ACTGGAACATGAAG 727
                                                                                                                                                                                                                                                                                                                                      AAV08201 standard; DNA; 21
                                                                                                                                                                                                                                                          4 ACTGGAACATGAAG 17
                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
                                                                                                                                                                                                14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Nathans J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 1998-495375/42.
                                                                                                                                                                                Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   27-FEB-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                    27-JAN-1999
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Lupski JR,
Sun H;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       03-SEP-1998.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Synthetic.
                                                                                                                                                                                                                                                                                                                                                                     AAV08201;
                                                                                                                                                                 Query Match
                                                                                                                                                                                                                                                                                                           RESULT 1291
                                                                                                                                                                                                Matches
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Gaps

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0.8%; Score 14; DB 1; Length 21; 100.0%; Pred. No. 9.8e+02; iive 0; Mismatches 0; Indels

14; Conservative

Matches

Best Local Similarity

Query Match

Human; heparanase; gene therapy; tumour; inflammation; autoimmunity; heparin-binding growth factor; cytokine; neurodegenerative plaque; wound healing; infection; burn; amplogenesis; restenosis; atherosclerosis; inflammation; neurodegenerative disease; Gerstmann-Straussler Syndrome; Creutzfeldt-Jakob disease; PCR primer; ss.

WO200052178-A1. Homo sapiens.

08-SEP-2000.

PCR primer hpl-629 used to amplify human cDNA encoding heparanase.

(first entry)

15-JAN-2001

AAA75055;

AAA75055 standard; DNA; 21

RESULT 1293

10017621-381.rng

AAA75055

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The specification describes a polypeptide having heparanase (hpa) activity. The recombinant protein is used as a modulator of heparin-binding growth factors, cellular responses to heparin-binding growth factors and cytokines, cellular responses to heparin-binding growth factors and cytokines, cell interaction with plasma lipoproteins, cellular susceptibility to viral, protozoal and bacterial infections or disintegration of neurodegenerative plaques. Heparanase may be useful for athersclerosis, inflammation, neurodegenerative diseases, and viral infections. Mammatian heparanase can be used to neutralize plasma heparin, and anti-heparanase antibodies may be applied for immunodetection and diagnosis of micrometastases, autoimmune lesions, and present parts of micrometastases, autoimmune lesions, and present parts which is amplify has cDNA, in the course of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New human polynucleotide useful for treating angiogenesis, restenosis, and inflammation.
                                                                                                                                                                                                                                                       Heparanase; hpa; modulator; heparin-binding growth factor; cellular response; cytokine; cell interaction; plasma lipoprotein; cellular susceptibility; infection; disintegration; neurodegenerative plaque; wound healing; angiogenesis; restenosis; atherselerosis; inflammation; neurodegenerative disease; neutralise; plasma heparin; micrometastasis; autoimmune lesion; renal failure; pcr primer; ss.
                                                                                                                                                                                                                     PCR primer used to amplify human heparanase cDNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 21 BP; 6 A; 8 C; 6 G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (INSI-) INSIGHT STRATEGY & MARKETING LTD.
(HADA-) HADASIT MEDICAL RES SERVICES & DEV.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Feinstein E;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Example 7; Page 30; 63pp; English.
                                                                                                                       BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   98WO-US017954.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   97US-00922170.
   717
                                8 AGGAGATCAGACTG 21
                                                                                                                     AAX35653 standard; DNA; 21
                                                                                                                                                                                       (first entry)
 704 AGGAGATCAGACTG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Pecker I, Vlodavsky I,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FRIEDMAN M M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 1999-302255/25.
                                                                                                                                                                                       09-JUL-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 31-AUG-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                             WO9911798-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    02-SEP-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   02-JUL-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 11-MAR-1999
                                                                                                                                                                                                                                                                                                                                                                                               Synthetic.
                                                                                                                                                       AAX35653;
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                                                                                                   AAX35653,
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The present PCR primer was used to amplify a human cDNA sequence, which encoded a protein with heparanase catalytic activity. The heparanase (hps) polynucleotide is useful in gene therapy, particularly in treating tumour, inflammation or autoimmunity. Particularly, the polynucleotide is useful in modulating the bioavailability of heparin-binding growth factors, cellular responses to heparin-binding growth factors (e.g. bFGF) and cytokines (e.g. interleukin (IL)-8), cell interaction with plasma and cytokines, cellular susceptibility to certain viral and some bacterial protozoa infections, or disintegration of neurodegenerative plaques. The polynucleotide is also useful in wound healing (e.g. thermal, chemical or radiation burns) and in the treatment of anglogenesis, restenosis, atherosclerosis, inflammation, neurodegenerative diseases (Gerstmann-Straussler Syndrome or Creutzfeldt-Jakob disease), and some
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New polynucleotides encoding a polypeptide having heparanase activity, useful in wound healing and in gene therapy, particularly in treating tumor, inflammation, autoimmunity, neurodegenerative diseases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
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0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 21 BP; 6 A; 8 C; 6 G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                      (INSI-) INSIGHT STRATEGY & MARKETING LTD.
(HADA-) HADASIT MEDICAL RES SERVICES & DEV.
(PRIE/) FRIEDMAN M M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 bacterial or protozoa infections
                                                                                                                                                                                                                                                                                                                                                                                                                          Feinstein E;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example 6; Page 53; 152pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAH28645 standard; DNA; 21 BP.
                                                                                                                                                                                                                                                                                           14-FEB-2000; 2000WO-US003542.
                                                                                                                                                                                                                                                                                                                            99US-00258892.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               273 TGCTGCTCCTGGGG 286
                                                                                                                                                                                                                                                                                                                                                                                                                         Vlodavsky I,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2000-579289/54.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Similarity
                                                                                                                                                                                                                                                                                                                         01-MAR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                         Pecker I,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               14
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1294
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AAH28645
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Gaps 0;

Length 21; 0; Indels

Score 14; DB 1; Le Pred. No. 9.8e+02;

0.00, 100.0%; Pre-

Query Match Best Local Similarity

Conservative

14;

Best Loca Matches

273 TGCTGCTCCTGGGG 286

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recrectereded 1

14

Mismatches

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The present invention relates to selection systems for the identification of cell proliferation genes based on functional analysis. A process is provided for the identification of a cell proliferation promoting cell proliferation promoting cell proliferation promoting acceptable isolation of genes involved in such activity, and the use of these genes for the diagnosis or treatment of a disease associated with excessive cell proliferation. The cell proliferation gene may be an oncogene, a dominant transforming gene, a tumour suppressor gene or a concept and the control of apoptosis. Antibodies, peptides and control of acceptable and the identified gene and/or its gene product for the treatment of cancer, arteriosclerosis, psoriasis, rheumatoid arthritis and restenosis cancer, arteriosclerosis, psoriasis, rheumatoid arthritis and restenosis cancer, arteriosclerosis, psoriasis, rheumatoid arthritis and restenosis cancer, arteriosclerosis, mutagenic agents termed perturbagens.

CC ancer, arteriosclerosis, and the gene(s) that allow escape from arrest are identified. The present sequence is that of PCR primer ovT201, which is homologous to a region within the PGK1 3' untranslated region.

CT he primer was used in an example from the invention in which the
                                                                                                                                                                                                                                                                       Identifying cell proliferation genes for treating diseases related to unregulated proliferation, by selecting revertant cell lines, analyzing their gene expression pattern and identifying differentially expressed genes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           determine the general efficacy of a screen for perturbagen molecules
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Cell proliferation; cellular target; viral growth; perturbagen; PCR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     pheromone response pathway of Saccharomyces cerevisiae was used to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 14; DB 1; Length 21;
Pred. No. 9.8e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 21 BP; 6 A; 1 C; 8 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0; Mismatches

 cerevisiae PGK1 PCR primer oVT201.

                                                                                                                                                                                                                                                                                                                                                                                                Example 4; Page 30; 42pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  100.08;
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97US-00812994.
                                            02-AUG-2001; 2001US-00921101.
                                                                                           99US-00252204.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AGCGTAAAGGATGG 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match 0.8
Best Local Similarity 100.
Matches 14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Saccharomyces cerevisiae.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         8 AGCGTAAAGGATGG
                                                                                                                                                                                                                                  WPI; 2002-328583/36.
                                                                                                                                          (ARCA-) ARCARIS INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  US2002132229-A1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            19-AUG-1996;
04-MAR-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     27-FEB-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                19-SEP-2002,
                                                                                           18-FEB-1999;
14-FEB-2002.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ABS57693;
                                                                                                                                                                                         Kamb CA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             RESULT 1296
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The present invention provides the protein, cDNA and genomic sequences of human interleukin-13 (IL13), and describes the single nucleotide bolymorphisms (SNRS) found within the gene, which is found on chromosome 5q31. IL13 is a pro-inflammatory cytokine thought to be involved in the pathogenesis of asthma and other immune and inflammatory diseases. The IL13 sequences and the SNPs identified can be used in drug screening, to determine an individual's susceptibility to disease, in forensic and paternity testing, and to identify treatments for cancer, immune and inflammatory diseases, including asthma and diseases characterised by
                                                                                                  interleukin-13; IL13; single nucleotide polymorphism; SNP; cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Novel polymucleotide comprising single nucleotide polymorphisms in human interleukin-13 gene is useful for studying expression and function of interleukin-13, as well as diagnosing and treating cancer, inflammatory,
                                                                                                                      inflammation, immune disorder, cytokine, asthma; chromosome 5q31; fibrosis, forensic, disease susceptibility; drug screening; PCR primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gene identification; cell proliferation; cancer; arteriosclerosis; psoriasis; rheumatoid arthritis; restenosis; gene therapy; cytostatic; antiarteriosclerotic; antipsoriatic; antiarthritic; antirheumatic; vasotropic; diagnosis; perturbagen; PGK1; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     fibrosis. The present sequence is an IL13 fragment PCR primer
                                                   Human interleukin-13 coding sequence fragment PCR primer #20.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0.8%; Score 14; DB 1; Length 21; 00.0%; Pred. No. 9.8e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 21 BP; 5 A; 7 C; 5 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Nandabalan K, Stephens JC;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               100.0%; Prec. ...
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ABL53717 standard; DNA; 21 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                            PHARM INC
                                                                                                                                                                                                                                                                                                                                                                27-SEP-2000; 2000WO-US026556
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       (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity 100.º
Matches 14; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PGK1 PCR primer oVT201.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           and immune disorders.
                                                                                                                                                                                                                                                                                                                                                                                                                                                         (GENA-) GENAISSANCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Denton RR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2001-343160/36
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                                                                                                                                                                                                                                                                       WO200123410-A2
                                                                                                                                                                                                                              Homo sapiens.
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         17-JUL-2001
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Chew A,

ABL53717;

RESULT 1295 ABL5371

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Gaps

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0; Indels

(BRIM) BRISTOL-MYERS SQUIBB CO

18-JAN-2002; 2002US-0350061P. 17-JAN-2003; 2003WO-US001981

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(ARCA-) ARCARIS INC.
           Poritz MA;
               WPI; 2003-138536/13
19-AUG-1997;
  06-NOV-1997;
   26-FEB-1999;
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           Kamb
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; 0.8%; Score 14; DB 1; Length 21; 100.0%; Pred. No. 9.8e+02; 0; Indels Sequence 21 BP; 6 A; 1 C; 8 G; 6 T; 0 U; 0 Other; 100.0%; Preq. ... Local Similarity 100. Les 14; Conservative Query Match Best Loc Matches

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Gaps

01-JAN-2004 ADD14266; RESULT 1297 ADD14266

Human src biomarker forward PCR primer SEQ ID NO:455.

Homo sapiens Synthetic

WO2003062395-A2

31-JUL-2003

This invention describes a novel method for identifying a cell

Call. The method comprises: (a) identifying within a number of virally
infected cells a cell that continues to proliferate; and (b) identifying
infected cells a cell that continues to proliferate; and (b) identifying
infected cells a cell that continues to proliferate a corresponding cell

Co proliferation gene or cellular target. The invention also describes a
method for identifying a perturbagen that inhibits viral growth. The cell

Co proliferation gene or cellular target. The invention also describes a
method for identifying a perturbagen that inhibits viral growth. The cell

Co proliferation gene or cellular target. The invention describes a
method for identifying a perturbagen that inhibits viral growth. The cell

Co unregulated cell proliferation, or for the development of antisense
capproaches and ribozymes. As the method involves positive selection,

i.e., selection for growth, rather than cessation of growth, it is easier

co identify and separate growing cells from growth arrested cells than to

co isolate non-transformed revertants. Since cultured tumour cell lines grow

vigorously in culture, the method can be performed in a time-efficient

co isolate non-transformed revertants. Since cultured tumour cell lines grow

vigorously in culture, the method can be performed in a time-efficient

co isolate non-transformed solonies can be identified, isolated, and analysed

cot the invention, as is the case in assays based on selection for non
transformed cells. This sequence represents a PCR primer used with the

primer represented in ABS57694 which is capable of amplifying the yeast

(Saccharomyces cerevisiae) PGKI 3'UTR which is used in a construct to Identifying cell proliferation gene involved in viral growth, comprises identifying cell that continues to proliferate within virally infected cells, and identifying corresponding cell proliferation gene in identified cell. Example 4; Page 30; 43pp; English. 97WO-US014514. 97US-00965477. 99US-00259155.

ADD14266 standard; DNA; 21 BP 8 AGCGTAAAGGATGG 21 6 AGCGTAAAGGATGG 19 (first entry) g à

predictor set; protein tyrosine kinase activity modulator; protein tyrosine kinase pathway; protein tyrosine kinase; cytostatic; gene therapy; drug sensitivity; genetic profile; cancer; human; PCR primer;

The present invention describes a predictor set comprising a plurality of polymucleotides or polypeptides whose expression pattern is predictive of the response of cells to treatment with a compound that modulates protein tyrosine kinase activity or members of the protein tyrosine kinase activity of cells, comprising obtaining a sample of cells, pathway. Also described: (1) predicting whether a compound is capable of modulate the activity of cells, comprising obtaining a sample of cells, determining whether the cells express a plurality of markers, and correlating the expression of the markers to the compound's ability to determining whether the cells; (2) a plurality of cells inness for identifying polymucleotides and polypeptides whose expression levels correlate with compound sensitivity or resistance of cells associated with a disease state; and (3) identifying polymucleotides and cells associated with a disease state, comprising subjecting the plurality of associated with a disease state, comprising subjecting the plurality of cell lines to one or more compounds, analysing the expression pattern of cells associated with a disease state by using the expression pattern of the microarray of polymucleotides and polypeptides that predict the sensitivity or resistance of cells associated with a disease state by using the expression pattern of the microarray. The polymucleotides and polypeptides are useful in predicting the activity of compounds that interact with protein tyrosine kinases and/or protein tyrosine kinase pathways. These may be used in determining drug sensitivity in patients to allow the development of individualized canner invention in the acceptance of ances of an ances of a patient response at a molecular level. The present invention is the acceptance of an expension patient prosent invention of present inventions. New polynucleotides and polypeptides for predicting the activity of compounds that interact with protein tyrosine kinases and/or protein sequence is used in the exemplification of the present invention. Sequence 21 BP; 6 A; 5 C; 6 G; 4 T; 0 U; 0 Other; Shaw P; Example 2; SEQ ID NO 455; 139pp; English. Lee FY, tyrosine kinase pathways. Huang F, Fairchild CR, WPI; 2003-636735/60. Query Match

Gaps . 0 / Match 0.8%; Score 14; DB 1; Length 21; Local Similarity 100.0%; Pred. No. 9.8e+02; nes 14; Conservative 0; Mismatches 0; Indels Matches

.. 0

245 GCAGTGACCCTGGA 258 20 GCAGTGACCCTGGA

> à g

ACH00878 standard; DNA; 21 (first entry) 12-FEB-2004 ACH00878;

BP

RESULT 1298

ACH0087

L monocytogenes CtsR protein fragment coding sequence.

CtsR; glycine-rich region; stress tolerance; virulence; motility; fermentation; vaccine; antibacterial; Class III stress gene regulator; gene; ds.

Listeria monocytogenes.

Location/Qualifiers

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WPI; 2003-874843/81.
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                                                                                                                                            (ILAN/)
                                                                                                                                                              (VLOD/)
(YACO/)
(PECK/)
                                                                                                                                                                                                                            (FEIN/)
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HANDER STANDER STANDER
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The present invention relates to a nucleic acid molecule which encodes an altered Class III stress gene regulator (CtsR) protein of a Gram-positive becterium. Such an altered CtsR protein has an alteration in the conserved glycine-rich region that corresponds to amino acid positions 61-64 of the normal CtsR protein. The alteration confers to the altered CtsR protein increased stress tolerance, reduced virulence or reduced mobility of a Gram positive bacterium in which the altered CtsR protein is expressed as sole CtsR protein. The coding sequence encoding an altered CtsR protein. The coding sequence encoding an attered CtsR protein is useful in a fermentation process, as a probiotic or as a live oral vaccine. The present sequence is a coding sequence for the fragment of the Listeria monocytogenes CtsR gene which encodes the
                                                                                                                                                                                                                                                                                                                                                                                                                                  New nucleic acid molecule comprising a nucleotide sequence encoding an altered Class III stress gene regulator (CtsR) protein of a Gram-positive bacterium, useful in a fermentation process, as a probiotic or as a live oral vaccine.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Wound healing; heparanase; ulcer; burn; laceration; surgical incision; necrosis; pressure wound; diabetic ulcer; angiogenesis; human; therapy; RACE; rapid amplification of cDNA end; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
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100.0%; Pred. No. 9.8e+02;
cive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human hpa specific antisense RACE PCR primer, hpl-629.
                     /*tag= a
/product= "CtsR glycine-rich region"
                                                                                                                                                                                                                                                                                                                                       Karatzas KA, Bennik MHJ, Abee T, Kleerebezem M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 21 BP; 4 A; 2 C; 9 G; 6 T; 0 U; 0 Other;
                                                                                      /note= "no start or stop codon"
                                                                                                                                                                                                                                                                                             (WAGE-) WAGENINGEN CENT FOOD SCI.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example; Fig 1; 46pp; English.
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                                                                                                                                                                                                                                                  11-MAR-2002; 2002EP-00075946.
                                                                                                                                                                                                               11-MAR-2003; 2003WO-NL000178
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         230 GTGGTGGTGGTGGC 243
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Best Local Similarity 100.
Watches 14; Conservative
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                                                                'partial
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        glycine-rich region
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The present invention relates to methods and compositions for inducing and/or accelerating wound healing via the catalytic activity of heparanase. The invention is used to induce or accelerate a healing process, particularly of an ulcer, burn, laceration, surgical incision, necrosis, pressure wound, diabetic ulcer and to induce or accelerate angiogenesis. The present sequence is human hpa specific antisense RACE
                                                                                                                                                                                                                                                                                                                                                                                               Composition for treating a wound comprising recombinant heparanase is useful to induce or accelerate wound healing and induce or accelerate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ss; chemical modification; methylation; array; CpG island; suppressor; p16; human; H69; H1618.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human DNA probe used to immobilise CpG methylated DNA SeqID 1096.
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0
                                                                                                                                                                                                                                                                                              Feinstein E;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 14; DB 1; Length 21;
Pred. No. 9.8e+02;
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                                                                                                                                                                                                                                                                                                Pecker I,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 21 BP; 6 A; 8 C; 6 G; 1 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                              Yacoby-Zeevi O,
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               01-WAR-1999, 99US-00258892.
06-FEB-2001, 2001US-00776874.
05-SEP-2001; 2001WO-IL000830.
19-NOV-2001; 2001US-00988113.
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98WO-US017954
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                                                                                                                                                                                                                                                                                                   Vlodavsky I,
                                                                                                                                                                     VLODAVSKY I.
YACOBY-ZEEVI (
PECKER I.
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MINNA J D.
LUEBKE K J.
BALOG R P.
                                                                                                                                                                                                                                                    FEINSTEIN E.
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of chemical modification of DNA involves obtaining sample of DNA to be analyzed, treating DNA with chemical reagents that result in different base sequences, and determining sequence of resulting DNA. Analysis

Example 1; SEQ ID NO 1096; 210pp; English.

This invention relates to a novel method for analysing chemically modified macromolecules. Specifically, it refers to a high throughput method for the parallel analysis of many potential sites of chemical modification (e.g. methylation) in DNA. The present invention describes treating the DNA with one or more chemical reagents that result in different base sequences depending upon the presence or absence of the modification of interest. Accordingly, a device comprising an array of probes is provided to hybridise with and select the altered DNA sequences that comprise the modifications of interest such as a CpG island. In particular, this invention refers to analysing the methylation pattern of human lung tumour cell lines H69 and H1618. This oligonucleotide sequence is a human DNA probe used to immobilise CpG methylated DNA of the

Sequence 21 BP; 4 A; 11 C; 3 G; 3 T; 0 U; 0 Other;

Gaps . 0.8%; Score 14; DB 1; Length 21; 100.0%; Pred. No. 9.8e+02; rative 0; Mismatches 0; Indels 0.8%; 14; Conservative Sest Local Similarity Query Match

555 CCTCAGCCGCCGCC 568 cercadededece 21

RESULT 1301 ADJ14006

ADJ14006 standard; DNA; 21 BP.

20-MAY-2004 (first entry) THE STATE OF THE S

probe; ss; chemical modification; methylation; array; CpG island; tumour suppressor; p16; human; H69; H1618.

Human DNA probe used to immobilise CpG methylated DNA SegID 1133.

Homo sapiens

US2003152950-A1

14-AUG-2003.

27-JUN-2002; 2002US-00184085.

27-JUN-2001; 2001US-0301370P

(GARN/) GARNER H R. (MINN/)

MINNA J D. LUEBKE K J. BALOG R P. (BALO/) (LUEB/)

Luebke KJ, Sarner HR, Minna JD,

Balog RP;

WPI; 2003-874843/81.

Analysis of chemical modification of DNA involves obtaining sample of DNA to be analyzed, treating DNA with chemical reagents that result in different base sequences, and determining sequence of resulting DNA.

Example 1; SEQ ID NO 1133; 210pp; English.

This invention relates to a novel method for analysing chemically modified macromolecules. Specifically, it refers to a high throughput

method for the parallel analysis of many potential sites of chemical modification (e.g. methylation) in DNA. The present invention describes treating the DNA with one or more chemical reagents that result in different base sequences depending upon the presence or absence of the modification of interest. Accordingly, a device comprising an array of probes is provided to hybridise with and select the altered DNA sequences that comprise the modifications of interest such as a CpG island. In particular, this invention refers to analysing the methylation pattern of a region of the promoter for the tumour suppressor game pl6 from two human lung tumour cell lines H69 and H1618. This oligomocleotide sequence is a human DNA probe used to immobilise CpG methylated DNA of the

Sequence 21 BP; 4 A; 11 C; 3 G; 3 T; 0 U; 0 Other;

Gaps ·, Score 14; DB 1; Length 21; Pred. No. 9.8e+02; 0; Indels 0.8%; scc... 100.0%; Pred. No. ... 0; Mismatches Query Match 0.8 Best Local Similarity 100. Matches 14; Conservative

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ADL16386/

ADL16386 standard; DNA; 21 BP.

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ADL16386;

06-MAY-2004 (first entry)

Human heparanase 5' RACE primer hpl-629.

Human; ss; heparanase; PCR; primer; heparanase-dependent cancer; cancer; autoimmune reaction; inflammation.

Homo sapiens.

US2003236215-A1.

25-DEC-2003.

09-JUN-2003; 2003US-00456573

98WO-US017954. 31-AUG-1998; 01-MAR-1999;

99US-00258892. 99US-00435739. 08-NOV-1999; (INSI-) INSIGHT STRATEGY & MARKETING LTD. (HADA-) HADASIT MEDICAL RES SERVICES & DEV.

Pecker I, Vlodavsky I, Feinstein E;

WPI; 2004-070610/07.

New antisense oligonucleotide hybridizable with a polynucleotide encoding a polypeptide with heparanase activity, useful for treating diseases such as cancer and autoimmune disorders.

Example 6; SEQ ID NO 17; 108pp; English.

The invention relates to an antisense oligonucleotide (ASO) comprising a polynucleotide or a polynucleotide analogue of at least 10 bases being hybridisable in vivo, under physiological conditions, with a portion of a polynucleotide strand encoding a polypetide having heparanase catalytic activity. Also included are a method of in vivo downregulating heparanase activity (comprising administering the ASO in vivo), a method of treating a subject suffering from a pathological condition (characterised by heparanase activity, comprising ASO to the subject), a pharmaceutical composition comprising the ASO and a carrier, an antisense nucleic acid construct (comprising a promoter sequence and a polynucleotide sequence directing the synthesis of an

Example 6; SEQ ID NO 17; 106pp; English.

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under physiological conditions, with a polynucleotide strand encoding a polypeptide having heparanase catalytic activity), a method of in vivo downregulating heparanase activity (comprising administering in vivo the antisense nucleic acid construct), a pharmaceutical composition comprising the antisense nucleic acid construct and a carrier, and an attisense obligonucleotide comprising a polynucleotide or a polynucleotide antisense solidonucleotide comprising a polynucleotide or a polynucleotide comprising at least 10 bases being hybridisable in vivo, under physiological conditions, with a portion of a polynucleotide strand being characterised by forming at least a portion of an untranslated region (UTR) for a polynucleotide strand encoding a polypeptide having chearanase catalytic activity. The methods and compositions of the present invention are useful for the prevention and/or treatment of diseases or conditions associated with aberranase activity, such as heparanase-dependent cancer, cancer, autoimmune reaction and continuous inflammation. The present sequence is a PCR primer used in the isolation
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antisense RNA sequence of at least 10 bases being hybridisable in vivo
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                                                                                                                                                                                                                                                                                                       Sequence 21 BP; 6 A; 8 C; 6 G; 1 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human hpa DNA amplifying PCR primer, hpl-629.
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06-FEB-2001; 2001US-00776874.
19-NOV-2001; 2001US-00988113.
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Chajek-Shaul T, Goldshmidt
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CHAJEK-SHAUL T.
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VLODAVSKY I.
METZGER S.
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(METZ/)
(PECK/)
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New transgenic non-human animal expressing heparinase, useful as models for human disease, such as cancers, viral infection, neurodegenerative diseases, restenosis, atherosclerosis and pulmonary disorders.

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The present invention relates to a primer set and method for mapping at least the variation/mutation part of a plant gene using a gene polymorphism marker. A mutation site of the plant gene is mapped by utilizing a genetic polymorphism marker as follows: (a) genomic DNA is prepared from a plant homozygously having a mutation to be an object of the mapping; (b) A forward primer 1 contealing a base corresponding to the gene polymorphic maker of none ecotype plant, a forward primer 2 containing a base corresponding to the genetic polymorphism of the other ecotype plant and a reverse primer 3 based on the base sequence common with both the ecotype plants are prepared; (c) two kinds of oligonucleotides emitting fluorescence of different colors when the genetic polymorphism marker is detected are prepared; (d) an
                       The present invention relates to a transgenic non-human animal whose genome comprises an exogenous polynucleotide sequence, including a promoter active in tissues of the non-human, a region encoding a human heparanase, where the promoter and the region encoding human heparanase are expressed in at least a portion of the cells of the non-human animal. The methods and compositions of the present invention are useful for the production of transgenic animals expressing heparanase, be used as models for human diseases such as cancers, viral infection, restenosis, neurodegenerative diseases, atherosclerosis and pulmonary disorders. The present sequence is human hpa DNA amplifying PCR primer
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                                                                                                                                                                                                                                                                                                          Score 14; DB 1; Length 21;
Pred. No. 9.8e+02;
                                                                                                                                                                                                                                                                                                                                               0; Indels
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                                                                                                                                                                                                                                                                       Sequence 21 BP; 6 A; 8 C; 6 G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                                                                      used in the exemplification of the invention.
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Best Local Similarity 100.0%; Pred. No. 9.8
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                                                                                                                                                                                                                                                                                                                                                                                        273 TGCTGCTCCTGGGG 286
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MATSUI M.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ss; diagnosis; non-blood disease; solid tumor; gene expression; peripheral blood mononuclear cell; renal cell carcinoma; prostate cancer; head/neck cancer; differential expression; probe.
amplification reaction of the genomic DNA is carried out in the presence of the primers 1, 2 and 3 and the two kinds of the oligonucleotides; (e) the fluorescence intensity emitted from the resultant reactional product is detected and (f) the position on the genome of the mutation site is determined from the results of detection. The present sequence is a primer, used to illustrate the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The invention relate to a method of diagnosing (M1) non-blood disease such as solid tumor by providing peripheral blood sample of human having non-blood disease, and comparing an expression profile of specific genes in the peripheral blood sample to reference expression profile of the genes, where each of the genes is differentially expressed in peripheral blood mononuclear cells (PBMCs) of patients having the disease as blood mononuclear cells (PBMCs) of patients having the disease as compared to PBMCs of normal humans. The method is useful for diagnosing non-blood disease such as solid tumor. The solid tumor is chosen from renal cell carcinoma (RCC), prostate cancer and head/neck cancer. The peripheral blood sample comprises enriched PBMCs. The peripheral blood sample (claimed). (M1) is useful for identifying
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Diagnosing non-blood disease such as solid tumor, involves comparing differential expression profile of specific genes in peripheral blood sample of subject with reference expression profile of specific genes.
                                                                                                                                                                                             Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                Renal cell carcinoma differentially expressed gene probe #2090.
                                                                                                                                                       Score 14; DB 1; Length 21; Pred. No. 9.8e+02;
                                                                                                                                                                                         0; Indels
                                                                                                                     Sequence 21 BP; 4 A; 7 C; 3 G; 7 T; 0 U; 0 Other;
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03-APR-2003; 2003US-0459782P.
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                                                                                                                                                           0.8%;
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BURCZYNSKI M E.
TREPICCHIO W L.
DORNER A.
STOVER J A.
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                                                                                                                                                                         Local Similarity
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(BURC/)
(TREP/)
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                                                                                                                                                                                                                                                                                                            RESULT 1305
ADP15685
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genes that are differentially expressed in peripheral blood samples isolated at different stages of progression, development or treatment of RCC and/or other solid tumors. This sequence corresponds to a probe to detect a gene that is differentially expressed and detected by the method
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                                                                                                                                  Gaps
                                                                                                                                                                                                                                                                                                                                     Rat ICAM hammerhead ribozyme target sequence (nt. position 510).
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                                                                                                    0.8%; Score 14; DB 1; Length 25; 77.3%; Pred. No. 1.1e+03;
                                                                                                                              5; Indels
                                                                           Sequence 25 BP; 6 A; 6 C; 4 G; 9 T; 0 U; 0 Other;
                                                                                                                              0; Mismatches
                                                                                                                                                      1691 TCCCTGCTTACTCTCTGCCTAC 1712
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940S-0029123.
940S-00292620.
940S-0029320.
940S-00300000.
940S-00300000.
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94US-00222795.
94US-00224483.
94US-00227958.
94US-00227958.
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94US-00357577.
94US-00363233.
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94US-00316771
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                                                                                                                                                                                                                                                                                                              (first entry)
                                                                                                                           Matches 17; Conservative
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                                                                                                                Best Local Similarity
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15-APR-1994;
15-APR-1994;
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28-SEP-1994;
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Claim 1; Page 76; 128pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                           Matches
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                                                                                                                                                                                                                                                                                                                     The present sequence represents a preferred target sequence for an enzymatic nucleic acid (i.e. a ribozyme) which cleaves ICAM-1 mRNA at the nucleotide base position indicated in the DB line. Regions of the mRNA that do not form secondary folding structures and that contain potential hammerhead and hairpin ribozyme cleavage sites were identified by computer analysis. Ribozymes directed against these mRNA sequences were designed and synthesised with modifications that improve their nuclease resistance. The ribozymes cleave the ICAM-1 target sequences and thereby inhibit ICAM-1 expression, making them useful for reducing transplant rejection and alleviating symptoms in patients with rheumatoid arthritis, correct PI field.)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New enzymatic nucleic acid molecules - cleave RNA produced by e.g. c-myb, for treating restenosis or cancer.
                                                                            o DT, Chowrira B, Direnzo A, Draper KG, Dudycz LW;
Karpeisky A, Kisich K, Matulic-Adamic J, Mcswiggen JA;
Pavco P, Beigleman L, Sullivan SM, Sweedler D, Thompson JD;
Usman N, Wincott FE, Woolf T;
                                                                                                                                                                                                                            having modified bases and methods for producing them - for use
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smooth muscle cell; hyperproliferation; restenosis; cancer; c-myb;
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                                                                                                                                                                                                                                                 in inhibiting disease related genes.
                                                                                                                                                                                                                                                                                          Claim 2; Page 201; 407pp; English.
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95US-00373124.
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  95US-00380734.
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                                          (RIBO-) RIBOZYME PHARM INC.
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Matches 12; Conservative
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                                                                                     Stinchcomb DT,
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    30-JAN-1995;
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Modak A,
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enzymatic nucleic acid, especially a hammerhead ribozyme, which cleaves the human c-myb sequence at the base position indicated in the descriptor line. The c-myb sequence was screened for optimal ribozyme target sites using a computer folding algorithm, and regions of the mRNA which did not form secondary folding structures and contained potential ribozyme cleavage sites were identified. Ribozymes were synthesised and their activities optimised by either varying the length of the binding arms or by modification to prevent degradation by nucleases. The ribozymes cleave the c-myb sequence and can be used to prevent smooth muscle cell
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The present sequence represents the preferred target sequence for an enzymatic nucleic acid, especially a hammerhead ribozyme, which cleaves the human c-myb sequence at the base position indicated in the descriptor line. The c-myb sequence was screened for optimal ribozyme target sites using a computer folding algorithm, and regions of the mRNA which did not form secondary folding structures and contained potential ribozyme cleavage sites were identified. Ribozymes were synthesised and their activities optimised by either varying the length of the binding arms or by modification to prevent degradation by nucleases. The ribozymes cleave the c-myb sequence and can be used to prevent smooth muscle cell.
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                                                                                                                                                                                                                                                                                                                                                              hyperproliferation in restenosis, especially after coronary angioplasty, and in cancers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Enzymatic nucleic acid; hammerhead; ribozyme; cleavage; human; smooth muscle cell; hyperproliferation; restenosis; cancer; c-myb;
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88.2%; Pred. No. 9e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 17 BP; 1 A; 3 C; 3 G; 0 T; 10 U; 0 Other;
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0; Mismatches
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95US-00373124.
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10017621-381.rng

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Sequence 17 BP; 2 A; 5 C; 5 G; 5 T; 0 U; 0 Other;
                                                    Query Match 0.8 Best Local Similarity 88.2 Matches 15; Conservative
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                                                                                                                                                                                                                                                                                                      Probe; interleukin-6 receptor; IL-6R; cytokine; cellular proliferation; transmembrane glycoprotein receptor; signal transducer; gp130; inhibitor; IL-6; cancer; renal cell carcinoma; autoimmune disease; viral infection;
restenosis, especially after coronary angioplasty,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Oligo:nucleotide(s) complementary to interleukin-6 receptor mRNA - for treating proliferative diseases, e.g. cancer, auto-immune diseases or viral infections.
                                                                                         Gaps
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                                                             Score 13.8; DB 1; Length 17;
Pred. No. 9e+02;
0; Mismatches 2; Indels
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/note= "optionally phosphorothioated"
                                      0 T; 10 U; 0 Other;
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                                      Sequence 17 BP; 1 A; 3 C; 3 G;
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95US-00486408.
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                                                           Query Match 0.8°
Best Local Similarity 88.2°
Matches 15, Conservative
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 hyperproliferation
              and in cancers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The present invention describes nucleic acid molecules which modulate the synthesis, expression and/or stability of a mRNA encoding 1 or more receptors of vascular endothelial growth factor (WGCF). A parient (preferably human) having a condition associated with the level of the fms-like tyrosine kinase 1 (ftt-1), kinase insert domain containing receptor (KDR) and/or foetal liver kinase 1 (flk-1) (e.g. tumour rangiogenesis, coular diseases, psociasis and rheumatoid arthritis) can be treated by administering the nucleic acid molecule or the expression vector to the patient. AAX67275 to AAX75752 represent specific examples of nucleic acid molecules from the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Vascular endothelial growth factor receptor; VEGF receptor; flt-1; flk-1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       KDR; hammerhead ribozyme; hairpin ribozyme; cleavage; tumour angiogenesis; psoriasis; rheumatoid arthritis; ocular disease; fme-like tyrosine kinase 1; kinase insert domain containing receptor; foetal liver kinase 1; se
                                                            Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Nucleic acid molecule modulating VEGF receptor(s) gene expression or stability - useful for treating e.g. tumour angiogenesis, psoriasis, rheumatoid arthritis, etc., in a human patient.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
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Score 13.8; DB 1; Length 17; Pred. No. 9e+02;
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54.7%; Pred. No. 9e+02;
ive 4; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human KDR VEGF receptor hammerhead ribozyme substrate #484.
                                                         Indels
                                                      2;
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                                                      Mismatches
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  0.8%;
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                                                                                                                                                                1 GGTGGACACCTCGTTCT
                                                                                                            1596 GGTGGACACCGAGTTCT
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                                                                                                                                                                                                                                                                                                    AAX71472 standard; RNA; 17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Pavco P, Mcswiggen J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 1997-259017/23.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CHIR ) CHIRON CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Local Similarity
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Gaps

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Indels

7

Mismatches

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The present invention describes enzymatic nucleic acid molecules with RNA cleaving activity, which specifically cleave RNA encoded by an aryl widrocarbon molear transporter (ARNT) gene, an integrin subunit beta 3 gene, an integrin alpha 6 subunit gene, or a Tie-2 gene. AAA16775 to AAA17167 and AAA17681 to AAA17682 to AAA18385 and AAA16775 to AAA19154 represent their corresponding target sequences; AAA17685 to AAA18385 and AAA19087 to AAA19154 represent ribozyme sequences for Tie-2, and AAA18385 to AAA19086 and AAA19155 to AAA19222 represent their corresponding target sequences; AAA19223 to AAA17685 to AAA18385 to AAA19086 to AAA19222 represent their corresponding target sequences; AAA21689 to AAA21681 represent their corresponding target sequences; AAA21689 represent their corresponding target sequences; AAA21689 to AAA21681 represent their corresponding target sequences; AAA21689 to AAA21683 represent their corresponding target sequences; cto integrin subunit beta 3, and AAA22476 to AAA23362, AAA23343 to the invention are used for modulating the synthesis, expression and/or stability of an mRNA encoding angiogenic factor, especially ARNT, integrin subunit beta 3, integrin subunit alpha-6, or Tie-2. They are especially used to treat cancer, diabetic retinopathy, age related macular degeneration (ARND), inflammation, and arthritis, as well as neovascular glaucoma, myopic degeneration, psoriasis, verruca vulgaris, and other syndrome, Kippel-Trenaunay-Weber septence telinopathy, age related syndrome, syndrome, whose syndrome, osler-Weber-Rendu syndrome, and other syndrome and diseases related to the levels of ARNT, Tie-2, and the levels of ARNT, Tie-2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 expression and/or stability
                                                                                                                                                                                                                                                                  integrin alpha 6 subunit; integrin subunit beta 3; hairpin ribozyme; hammerhead ribozyme; angiogenic factor; cytostatic; antidiabetic; oppthalmologic, antimilammatory; antiatrathritic; antiposriatic; ARMD; dermatological; RNA cleavage; cancer; diabetic retinopathy; arthritis; age related macular degeneration; inflammation; neovascular glaucoma; myopic degeneration; psoriaais; vertuca vulgaris; angiofibroma; tuberous sclerosis; pot-wine stain; Sturge Weber syndrome; ss. Kippel-Trenaunay-Weber syndrome; Osler-Weber-Rendu syndrome; ss.
                                                                                                                                                                                                                                                     ARNT; TIE-2; angiogenesis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mcswiggen JA;
                                                                                                                                                                                                   Integrin subunit beta 3 substrate sequence SEQ ID NO:6482.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Novel ribozymes for modulating the synthesis, of an mRNA encoding an angiogenic factors.
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                                                                                                                                                                                                                                                  aryl hydrocarbon nuclear transport;
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                                                  AAA23256 standard; RNA; 17 BP.
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                                                                                                                                                      (first entry)
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                                                                                                                                                      19-JUN-2000
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                                                                                                 AAA23256;
RESULT 1311
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DB 1; Length 17; Integrin subunit alpha-6, or integrin subunit beta-3 Score 13.8; DB Pred. No. 9e+02; 0.8%;

Query Match Best Local Similarity

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Human; c-raf; A-raf; B-raf; hammerhead ribozyme; hairpin ribozyme; target; substrate; catalyst; modulation; expression; Raf gene; delivery; screening; identification; synthesis; deprotection; purification; cancer; inflammation; psoriasis; non-hepatic ascites; infection; genetic drift; restenosis; rheumatoid arthritis; ss.
                                                                                                                                                                                                                                                                                                                                                                Bellon L;
                                                                                                                                                                                                                                                                                                                                                                          Burgin A;
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                                                                                                                                                                                                                                                                                                                                                                          Karpeisky A,
Sweedler D;
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                                                                                                                              Human A-Raf substrate position 1538.
                824
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97US-0051321P.
97US-0061324P.
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97US-0051718P
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97US-0068212P
                                                                                                                                                                                                                                                                                                                                                (RIBO-) RIBOZYME PHARM INC.
                 808 ATTATCCACACGGAGAA
                                                                            AAV92551 standard; RNA; 17
                                17 ATTATCCAAACGGAGCA
                                                                                                             (first entry)
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J, Workman CT,
15; Conservative
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09-JUN-1997;
03-JUL-1997;
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05-NOV-1997
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                                                                                                                                                                                                                                                                                                                                                                 Jarvis T,
                                                                                             AAV92551;
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                                                           RESULT 1312
Matches
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cleave target nucleic acid, particularly for treating systemic diseases caused by specific RNA, e.g. cancer, inflammation, psoriasis, non-hepatic ascites and infection. They may also be used to detect genetic drift and mutations in diseased cells and to determine c-raf RNA. Specifically NACs with RNA-cleaving activity that modulate expression of the Raf gene, are used to treat cancer, restenosis, psoriasis or rheumatoria artifuls, or generally any condition associated with the level of c-raf. Introduction of sugar/phosphate modifications increases stability against nuclease and activity. AAV90922 to AAV93877 represent NACs that can be used in the A method has been developed for the identification of a nucleic acid capable of modulating a process in a biological system. The method comprises: (a) introducing into the system a random library of nucleic acid catalysts (NAC) having a substrate binding domain (SBD), comprising a random sequence, and a catalytic domain (CD); and (b) identifying NAC in systems where modulation has occurred and/or determining the sequence of at least part of the SBDs in such systems. Nucleic acid molecules with endomuclease activity and catalytic activity, from the present invention, are used to modulate gene expression in plant and mammalian cells and to

Identifying new catalytic nucleic acid that modulates selected processes - especially ribozymes that cleave Raf RNA for treating cancer, restenosis, and also new ribozymes and modified nucleoside triphosphates

Beaudry A,

WPI; 1999-009494/01.

Thompson J,

used as antiviral agents and synthons.

Claim 177; Page 160; 259pp; English.

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                                                                 AAA72376;
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                RESULT 1314
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                            AAA72376/
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           A method has been developed for detecting the presence or absence of a single nucleotide polymorphism (SNP) allele in a genomic sample. The method comprises preparing a reduced complexity genome (RCG) from the genomic sample and analysing the RCG for the presence or absence of a SNP allele. The method can be used to characterise a tumour, to generate a genomic pattern for an individual genome or to generate a genomic classification code for a genome. The method can be used to assess whether a subject is at risk for developing a disease or to identify a set of SNP alleles associated with a disease. The method can also be used to perform linkage analysis. AAA35944 to AAA35947 represent sequences used in the exemplification of the present invention. AAA35948 to
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Detection of single nucleotide polymorphisms in genomes by preparation and analysis of reduced complexity genomes, useful for genotyping, fingerprinting and determining allele frequency of SNPs.
                                                                                                                                                                                                                                                                                                    Human; single nucleotide polymorphism; SNP; genotyping; DNA analysis; allels specific ollogonucleotide; ASO; reduced complexity genome; RCG; genomic classification; identification; DNA fingerprinting; tumour characterisation; hybridisation; SN.
                                                                               Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
method, specifically for modulating the expression of a Raf gene
                                                                                                                                                                                                                                                                             Human genomic SNP allele specific oligonucleotide SEQ ID NO:560.
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                                                   0.8%; Score 13.8; DB 1; Length 17; 58.8%; Pred. No. 9e+02; ative 5; Mismatches 2; Indels
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88.2%; Pred. No. 9e+02;
iive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAA36632 represent nucleotide sequences containing SNPs
                           Sequence 17 BP; 1 A; 3 C; 7 G; 0 T; 6 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Charest A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (MASI ) MASSACHUSETTS INST TECHNOLOGY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Housman DE,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Disclosure, Page 69; 111pp; English
                                                                                                      1030 GCTGACTTTGGCCTGGC 1046
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1112 CTGACATCCTGCTTGGG 1128
                                                                                                                      1 GGUGACUUUGGCUUGGC 17
                                                                                                                                                                                               AAA36495 standard; DNA; 17 BP
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                                                              Best Local Similarity 58.6
Matches 10; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2000-293181/25.
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Matches 15; Conserv
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                                                                                                                                                                                                                         AAA36495;
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II type 2 receptor (AT2 receptor) gene in vascular tissue. The invention also relates to a method for the production of transgenic animal of the invention, comprising inserting the AT2 receptor gene into pluripotent cells of the animal, implanting and bringing to term to give transgenic animals whose descendents will also express the AT2 receptor gene. The transgenic animal is a model system for the study of the vascular function and blood pressure regulatory function of the AT2 receptor in vivo or in vitro. It may also be used to study the competitive activity of AT1 and AT2 receptors. Sequences AAA72375-A72376 represent PCR primers represents a mouse AT2 receptor. PCR primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The invention relates to transgenic animals which express the angiotensin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Transgenic animals expressing angiotensin II2 receptor gene in vascular tissue used as a model for studying function and blood pressure regulatory activity of the receptor.
                                                                                                                                                                            Mouse angiotensin II type 2 receptor (AT2 receptor) PCR primer, AT2-R.
                                                                                                                                                                                                                                     Mouse angiotensin II type 2 receptor; AT2 receptor; vascular tissue; transgenic animal; blood pressure regulation; PCR primer; ss.
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88.2%; Pred. No. 9e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example 3; Page 9; 26pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      949 TACTGCCACCGCCAGAA 965
BP.
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AAA72376 standard; DNA; 17
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                                                                                                                 (first entry)
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                                                                                                                    19-DEC-2000
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Shannon ME;

Chen W,

Rank DR,

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New polypeptide, for raising antibodies that recognize hGDMLP-1 proteins, or as specific biomolecule capture probes for surface-enhanced laser
                                                                                                                                                                                                                                                                                                                                                     desorption ionization, comprises human myosin-like protein hGDMLP-1.
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2001WO-US000662.
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25-MAY-2001; 2001WO-US016981
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2001WO-US000666.
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30-JAN-2001; 2001WO-US000669
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05-FEB-2001; 2001US-0266860P.
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les 15; Conservative
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                                                                                                                                                                                                                                              (AEOM-) AEOMICA INC.
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30-JAN-2001;
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                                                    27-SEP-2000;
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                                                                                                                                                                                                                                                                                                                        The present invention relates to oligonucleotides based on nucleotide sequences obtained from both wild-type tubercle bacilli (wtTB) that are susceptible to a drug and mutant-type tubercle bacilli (mtTB) that are resistant to a drug. The drugs used in the present invention are rifampicin (RPP), streptomycin (SM), kanamycin (KM), isoniazid (INH) and thambutcl (BB). The rpoB gene is responsible for resistance to SR, the inhA gene is responsible for resistance to SM, the inhA gene is responsible for resistance to SM, the inhA gene is responsible for resistance to INH; the katG gene is responsible for resistance to INH; cand the embB gene is responsible for resistance to EB. The present invention also relates to nucleic acid probes having part of a nucleit sequence of tubercle bacilli (TB) responsible for drug resistance and primers used to generate the probes. The present sequence is an oligonucleotide of the present invention and be used to enable the differentiation of drug resistance and the determination of infection with tubercle bacilli
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                                                                                                                                                                                                                                             New oligonucleotides, nucleic acid probes and primers are useful for differentiating drug-resistance and determining infection with tubercle
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88.2%; Pred. No. 9e+02;
tive 0; Mismatches 2; Indels
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                                                                                                                                                                                                                                                                                                    Claim 16; Page 35; 114pp; English
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               Mycobacterium tuberculosis.
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                                                                                                                         03-AUG-1999;
                                          EP1076099-A2.
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ABN10018
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The present invention describes a human genome-derived myosin-like protein 1 (hGDMLP-1). The protein and polynucleotide sequences of hGDMLP-1 can be used as probes to detect, characterise and quantify nucleic acids can be used as probes to detect, characterise and quantify hGDMLP-1 nucleic acids in samples, as amplification substrates, to provide initial substrates for the recombinant engineering of hGDMLP-1 protein variants having desired phenotypic improvements, and for expressing the proteins. The hGDMLP-1 proteins or polypeptides may be used as immunogens to raise antibodies that specifically recognise hGDMLP-1 proteins, as standards in assays used to determine the concentration and/or amount specifically of hGDMLP proteins, as specific biomolecule capture probes for surface-enhanced laser description ionisation, as therapeutic supplement in patients having specific deficiency in hGDMLP-1 production, and in vaccines or for replacement therapy. The production, and in vaccines or for replacement therapy. The production and in vaccines or for seplacement therapy or diagnosing a disorder associated with the expression of hGDMLP-1, in particular heart and skeletal muscle disorders. hGDMLP-1 is localised to chromosome 22. The present sequence encounting and of hGDMLP-1, in particular heart hGDMLP-1 sequence in the exemplification of the present invention. N.B. The sequence data for this patent did not form part of the printed for the present sequence and sequence of the present invention.
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88.2%; Pred. No. 9e+02;
ive 0; Mismatches 2; Indels
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Human, genome-derived myosin-like protein 1; GDMLP-1; hGDMLP-1; heart; muscle; myosin; chromosome 22; gene therapy; vaccine; heart disease; skeletal muscle disorder; amplicon; screening; ss. 2000US-0236359P. 2000GB-00024263. 2001WO-US000661. 2001WO-US000662. 2001WO-US000663. 2001WO-US000664. 2001WO-US000665. 2001WO-US000666. 2001WO-US000667. 2001WO-US000668 2001WO-US000669 25-MAY-2001; 2001WO-US016981 2000US-0234687P 2001WO-US000670 05-FEB-2001; 2001US-0266860P WO200192524-A2 30-JAN-2001; 30-JAN-2001; 30-JAN-2001; 30-JAN-2001; 30-JAN-2001; 30-JAN-2001; 30-JAN-2001; 30-JAN-2001; Homo sapiens. 30-JAN-2001; 04-OCT-2000; 30-JAN-2001; 27-SEP-2000; 06-DEC-2001 21-SEP-2000

(AEOM-) AEOMICA INC.

Shannon ME; Chen W, Hanzel DK, Rank DR, Penn SG, WPI; 2002-179446/23. Ji Y,

New polypeptide, for raising antibodies that recognize hGDWLP-1 proteins, or as specific biomolecule capture probes for surface-enhanced laser desorption ionization, comprises human myosin-like protein hGDWLP-1.

Disclosure; SEQ ID NO 8045; 214pp; English.

The present invention describes a human genome-derived myosin-like protein 1 (hGDMLP-1). The protein and polynucleotide sequences of hGDMLP-1 can be used in gene therapy and vaccine production. The hGDMLP-1 collect acids can be used as probes to detect, characterise and quantify hGDMLP-1 nucleic acids in samples, as amplification substrates, to provide initial substrates for the recombinant engineering of hGDMLP-1 protein variants having desired phenotypic improvements, and for expressing the proteins. The hGDMLP-1 proteins or polypeptides may be used as immunogens to raise antibodies that specifically recognise hGDMLP-1 proteins, as specifically of hGDMLP-1 proteins, as specifically of hGDMLP-1 proteins, as specific biomolecule capture probes for surface-enhanced laser desorption ionisation, as therapeutic supplement in patients having specific deficiency in hGDMLP-1 production, and in vaccines or for replacement therapy. The production, and in vaccines or for replacement therapy. The production, and in vaccines or for replacement therapy. The production, and in vaccines or for replacement therapy. The colloquer associated with the expression of hGDMLP-1, in particular heart and skeletal muscle disorders. HGDMLP-1 is localised to chromosome 22.

The present sequence represents an oligomer used in the screening of the hGDMLP-1 sequence in the exemplification of the present invention. N.B. The sequence data for this patent did not form part of the present invention. N.B. The sequence data for this patent did not form part of the printed muscle dispersed in a patent did not form part of the printed muscle dispersed to the present invention. N.B. The sequence data for this patent did not form part of the printed manner of the present invention whit manner of the present invention in the manner of the present invention of the present invention when the manner of the present invention when the manner of the present invention the present invention the present invention of the printed the present invention the patent specification, but was obtained in electronic format directly from WIPO at ftp.wipo.int/pub/published_pct_sequence

Sequence 17 BP; 6 A; 2 C; 7 G; 2 T; 0 U; 0 Other;

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Gaps
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                      5
             9e+02;
0.8%; Score 13.8; D
88.2%; Pred. No. 9e+0
ative 0; Mismatches
                      Conservative
          Local Similarity
les 15; Conserv
 Query Match
                      Matches
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DB 1; Length 17;

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127 GATCGGATGAAGAGAT 143
                                                1 GAGCGGATGAAGCAGAT 17
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Human; genome-derived myosin-like protein 1; GDMLP-1; hGDMLP-1; heart; muscle; myosin; chromosome 22; gene therapy; vaccine; heart disease; skeletal muscle disorder; amplicon; screening; ss. Human GDMLP-1 17-mer scanning SEQ ID NO:5 sequence SEQ ID NO:6796. ABN06804 standard; DNA; 17 BP. 2000US-0234687P. 2000US-0236359P. 2000GB-00024263 2001WO-US000661 2001WO-US000662 2001WO-US000663 2001WO-US000664. 2001WO-US000665 2001WO-US000666. 2001WO-US000667 2001WO-US000668. 2001WO-US000669. 25-MAY-2001; 2001WO-US016981 2001WO-US000670 05-FEB-2001; 2001US-0266860P (first entry) WO200192524-A2. Homo sapiens. 30-JAN-2001; 30-JAN-2001; 30-JAN-2001; 30-JAN-2001; 30-JAN-2001; 30-JAN-2001; 30-JAN-2001; 30-JAN-2001; 30-JAN-2001; 27-SEP-2000; 29-MAY-2002 06-DEC-2001. ABN06804; RESULT 1318 ABN06804/

(AEOM-) AEOMICA INC.

Shannon ME; Chen W, Hanzel DK, Rank DR, Gu Y, Ji Y, Penn SG,

WPI; 2002-179446/23.

New polypeptide, for raising antibodies that recognize hGDMLP-1 proteins, or as specific biomolecule capture probes for surface-enhanced laser desorption ionization, comprises human myosin-like protein hGDMLP-1.

Disclosure; SEQ ID NO 6796; 214pp; English.

The present invention describes a human genome-derived myosin-like protein 1 (hGDMLP-1). The protein and polymucleotide sequences of hGDMLP-1 can be used in gene therapy and vaccine production. The hGDMLP-1 nucleic acids can be used as probes to detect, characterise and quantify hGDMLP-1 nucleic acids in samples, as amplification substrates, to provide initial substrates for the recombinant engineering of hGDMLP-1 protein variants having desired phenotypic improvements, and for expressing the proteins. The hGDMLP-1 proteins or polypeptides may be used as immunogens to raise antibodies that specifically recognise hGDMLP. I proteins, as standards in assays used to determine the concentration and/or amount specifically of hGDMLP proteins, as specific biomolecule capture probes for surface-enhanced laser description ionisation, as therapeutic supplement in patients having specific deficiency in hGDMLP-1 production, and in vaccines or for replacement therapy. The polynucleotide sequences encoding hGDMLP-1 may be used for diagnosing a disorder associated with the expression of hGDMLP-1, in particular heart and skeletal muscle disorders. hGDMLP-1 is localised to chromosome 2. The present sequence represents an oligomer used in the screening of the hGDMLP-1 sequence can the exemplification of the present invention. N.B. The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO

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27-SEP-2000;
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0.8%; Score 13.8; DB 1; Length 17;
Best Local Similarity, 88.2%; Pred. No. 9e+02;
Matches 15; Conservative 0; Mismatches 2: Indele
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                                         Sequence 17 BP; 0 A; 3 C; 11 G; 3 T; 0 U; 0 Other;
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  at ftp.wipo.int/pub/published_pct_sequence
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30-JAN-2001;
30-JAN-2001;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    26-MAY-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     21-SEP-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                04-OCT-2000;
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                                                                                                                                                                                                                                                                                                                                                       ABN01534;
                                                                                                                                                                                                                                                                       RESULT 1319
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and/or amount specifically of hGDMLP proteins, as specific biomolecule capture probes for surface-enhanced laser desorption ionisation, as preheargeutic supplement in patients having specific deficiency in hGDMLP-1 production, and in vaccines or for replacement therapy. The polymucleotide sequences encoding hGDMLP-1 may be used for diagnosing a disorder associated with the expression of hGDMLP-1, in particular heart and skeletal muscle disorders. hGDMLP-1 is localised to chromosome 22. The present sequence represents an oligomer used in the screening of the hGDMLP-1 sequence in the exemplification of the present invention. N.B. The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO at ftp.wipo.int/pub/published_pct_sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New polypeptide, for raising antibodies that recognize hGDMLP-1 proteins, or as specific biomolecule capture probes for surface-enhanced laser desorption ionization, comprises human myosin-like protein hGDMLP-1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human; genome-derived myosin-like protein 1; GDMLP-1; hGDMLP-1; heart; muscle; myosin; chromosome 22; gene therapy; vaccine; heart disease; skeletal muscle disorder; amplicon; screening; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human GDMLP-1 17-mer scanning SEQ ID NO:5 sequence SEQ ID NO:10664.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Shannon ME;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ·
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                0.8%; Score 13.8; DB 1; Length 17; 88.2%; Pred. No. 9e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        7
                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 17 BP; 3 A; 2 C; 9 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            9e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Disclosure; SEQ ID NO 10664; 214pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       986 AGCCCCAGAACCTGCTC 1002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABN10672 standard; DNA; 17 BP.
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2000US-0236359P.
2000GB-00024263.
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2001US-0266860P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            17 AGCCCCATCACCTGCTC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Local Similarity 88.2 ses 15, Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gu Y, Ji Y, Penn SG,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2002-179446/23.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WO200192524-A2
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10017621-381.rng

The present invention describes a human genome-derived myosin-like protein 1 (hGDMLP-1). The protein and polymucleotide sequences of hGDMLP-1 can be used in gene therapy and vaccine production. The hGDMLP-1 calcid caids can be used as probes to detect, characterise and quantify hGDMLP-1 nucleic acids in samples, as amplification substrates, to hGDMLP-1 nucleic acids in samples, as amplification substrates, to provide initial substrates for the recombinant engineering of hGDMLP-1 protein variants having desired phenotypic improvements, and for expressing the proteins. The hGDMLP-1 proteins or polypeptides may be used as immunogens to raise antibodies that specifically recognise hGDMLP-1 proteins, as standards in assays used to determine the concentration and/or amount specifically of hGDMLP-proteins, as specific biomolecule capture probes for surface-enhanced laser desorption ionisation, as therapeutic supplement in patients having specific deficiency in hGDMLP-1 production, and in vaccines or for replacement therapy. The polymucleotide sequences encoding hGDMLP-1 may be used for diagnosing a disorders associated with the expression of hGDMLP-1, in particular heart cand skeletal muscle disorders. hGDMLP-1 is localised to chromsome 2.

The present sequence represents an oligomer used in the screening of the hGDMLP-1 sequence and sceletal muscle disorders. hGDMLP-1 is localised to thromsome 2.

The sequence data for this patient did not form part of the printed concentration, hur was obtained in the exemplification of the present invention. N.B. ö specification, but was obtained in electronic format directly from WIPO at ftp.wipo.int/pub/published_pct_sequence Gaps .; 0 0.8%; Score 13.8; DB 1; Length 17; 88.2%; Pred. No. 9e+02; ative 0; Mismatches 2; Indels Sequence 17 BP; 4 A; 7 C; 6 G; 0 T; 0 U; 0 Other; Query Match
Best Local Similarity 88.2
Matches 15, Conservative

The present invention describes a human genome-derived myosin-like protein 1 (hGDMLP-1). The protein and polynucleotide sequences of hGDMLP-1 can be used in gene therapy and vaccine production. The hGDMLP-1 nucleic acids can be used as probes to detect, characterise and quantify hGDMLP-1 nucleic acids in samples, as amplification substrates, to provide initial substrates for the recombinant engineering of hGDMLP-1 protein variants having desired phenotypic improvements, and for expressing the proteins. The hGDMLP-1 proteins or polypeptides may be used as immunogens to raise antibodies that specifically recognise hGDMLP-1 proteins, as standards in assays used to determine the concentration

New polypeptide, for raising antibodies that recognize hGDMLP-1 proteins, or as specific biomolecule capture probes for surface-enhanced laser desorption ionization, comprises human myosin-like protein hGDMLP-1.

Disclosure; SEQ ID NO 6795; 214pp; English.

Chen W,

Rank DR,

Hanzel DK,

Gu Y, Ji Y, Penn SG, (AEOM-) AEOMICA INC.

WPI; 2002-179446/23.

and/or amount specifically of hGDMLP proteins, as specific biomolecule capture probes for surface-enhanced laser desorption ionisation, as therapeutic supplement in patients having specific deficiency in hGDMLP-1 production, and in vaccines or for replacement therapy. The polymolecotide sequences encoding hGDMLP-1 may be used for diagnosing a disorder associated with the expression of hGDMLP-1, in particular heart and skeletal muscle disorders. hGDMLP-1 is localised to chromosome 22.

The present sequence represents an oligomer used in the screening of the MGDMLP1 sequence in the exemplification of the present invention. N.B. The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from MIPO

ftp.wipo.int/pub/published_pct_sequence

> à qq

Human GDMLP-1 17-mer scanning SEQ ID NO:5 sequence SEQ ID NO:6795. ABN06803 standard; DNA; 17 BP 29-MAY-2002 (first entry) ABN06803; RESULT 1321 ABN06803,

Human, genome-derived myosin-like protein 1; GDMLP-1; hGDMLP-1; heart; muscle; myosin; chromosome 22; gene therapy; vaccine; heart disease; skeletal muscle disorder; amplicon; screening; ss.

Homo sapiens

06-DEC-2001

WO200192524-A2.

25-MAY-2001; 2001WO-US016981

2000US-0236359P. 2000GB-00024263. 2000US-0207456P 26-MAY-2000: 21-SEP-2000; 27-SEP-2000; 34-OCT-2000;

2001WO-US000661. 2001WO-US000662. 2001WO-US000663. 30-JAN-2001; 30-JAN-2001; 30-JAN-2001;

2001WO-US000664. 2001WO-US000665. 2001WO-US000666. 2001WO-US000670, 2001US-0266860P, 2001WO-US000667 2001WO-US000668 30-JAN-2001; 30-JAN-2001; 30-JAN-2001; 10-JAN-2001; 10-JAN-2001; 30-JAN-2001; 05-FEB-2001;

Gaps . 0.8%; Score 13.8; DB 1; Length 17; 38.2%; Pred. No. 9e+02; ve 0; Mismatches 2; Indels Sequence 17 BP; 1 A; 2 C; 11 G; 3 T; 0 U; 0 Other; 553 CCCCTCAGCCGCCGCT 569 ABQ63455 standard; DNA; 17 BP 88.2%; 17 ceccacaeccaecece 15; Conservative Best Local Similarity ABQ63455; Query Match RESULT 1322 ABQ63455/c Matches ò

; 0

Human, KTOMla; KTOMl; kidney tumour overexpressed membrane; cytostatic; gene therapy; cancer; kidney; liver; bone marrow; brain; heart; lung; kidney; colon; skeletal muscle; testis; uterus; placenta; probe; ss. Human KTOM1a portion (ABQ63232) probe # 168.

(first entry)

20-AUG-2002

Homo sapiens

WO200224750-A2.

28-MAR-2002.

21-SEP-2001; 2001WO-US029656.

27-SEP-2000; 2000US-0236359P. 04-CCT-2000; 2000GB-00024583 30-JAN-2001; 2001WO-US000661. 30-JAN-2001; 2001WO-US000662. 21-SEP-2000;

Randi AM;

Mclaughlin F,

Zhang J;

acids

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Expression of an Ets-related gene (ERG). (I) is useful for treating expression of an Ets-related gene (ERG). (I) is useful for treating conditions selected from cancer, lymphoma, Ewing's sarcoma, melanoma, tumour anglogenesis, diabetic retinopathy, macular degeneration, thorous anglofibroma of tuberous sclerosis, port-wine stains, Sturge Weber syndrome, Kippel-Trenaunay-Weber syndrome, Osteoporosis and wound healing. (I) is useful for syndrome, leukaemia, osteoporosis and wound healing. (I) is useful for treating a patient having a condition associated with the level of ERG, by contacting cells of the patient with (I) under conditions suitable for the treatment. The method comprises the use of one or more therapies under conditions suitable for the treatment or tumour conjunction with one or more of other therapies such as radiation or conjunction with one or more of other therapies such as radiation or conjunction with one or more of other therapies such as radiation or cell, by contacting (I) is useful for reducing ERG activity in a cell, by contacting (I) with RNA, in the presence of a divalent confuses related to the expression of ERG, and as diagnostic tool to examine genetic drift and mutations within diseased cells or to detect the presence of ERG RNA in a cell. (I) is useful for specifically carged enzymatic nucleic acid molecules which regulate expression of ERG, and carated PCR primers of the invention
                                                                                                                                                                                                                                                                                 Novel polynucleotide which down regulates expression of Ets-related gene, useful for treating cancer, diabetic retinopathy, macular degeneration, arthritis, psoriasis, verruca vulgaris and Sturge Weber syndrome.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     tumour angiogenesis; diabetic retinopathy; macular degeneration; neovascular glaucoma: myopic degeneration; arthritis; verruca vulgaris; angiofibroma of tuberous sclerosis; port-wine stain; wound healing; Sturge Weber syndrome; Kippel-Trenaunay-Weber syndrome; leukaemia; ss; osler-Weber-rendu syndrome; leukaemia; osteoporosis; DNAzyme; inozyme;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 17 BP; 1 A; 9 C; 4 G; 0 T; 3 U; 0 Other;
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70.6%; Pred. No. 9e+02;
                                                                                                                                                                               Mcswiggen JA,
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                                                                                                                                                                                                                                                                                                                                                                                               Claim 4; Page 82; 149pp; English
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16-MAY-2001; 2001WO-US015866.
                                                    16-MAY-2000; 2000US-00572021
                                                                                                                                                                                    Von Carlowitz I,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ABK18786 standard; RNA; 17
                                                                                                     (RIBO-) RIBOZYME PHARM INC. (GLAX ) GLAXO GROUP LTD.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity
                                                                                                                                                                                                                                        WPI; 2002-082995/11.
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                                                                                                                                                                                    Jarvis T,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                      w human kidney tumor overexpressed membrane (KTOM1) protein and nucleic ids encoding the protein, useful for treating subjects having defects KTOM1 which can manifest as cancer of the kidney, or as a disorder of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        KTOMI (kidney tumour overexpressed membrane) protein. The protein of the invention has cytostatic activity. The nucleotide may have a use in gene interapy. The KTOMI nucleic acids may be used to diagnose, treat or monitor a disease caused by altered expression of human KTOMI.

Compositions comprising the nucleic acids, proteins or antibodies may be used to treat subjects having defects in KTOMI which can manifest as cancer of the kidney, as well as a disorder of liver, bone marrow, brain, heart, lung, kidney, colon, skeletal muscle, testis, uterus and placenta function. The sequence represents a probe used in the invention to scan the nt 1-1001 portion of human KTOMIa (ABQ632312)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human; hammerhead ribozyme; cytostatic; antitumour; antidiabetic; ophthalmological; antiarthritic; antipsoriatic; virucide; osteopathic; vulnezry; cancer; lymphoma; Ewing's sarcoma; melanoma; psoriasis; tumour angiogenesis; diabetic retinopathy; macular degeneration; neovascular glaucoma; myopic degeneration; arthritis; verruca vulgaris; angiofibroma of tuberous solerosis; port.wine stain; wound healing; Sturge Weber syndrome; Kippel-Trenaunay-Weber syndrome; leukaemia; ss; osler-Weber-rendu syndrome, leukaemia; osteoporosis; DNAzyme; inozyme;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The invention relates to a novel isolated nucleic acid encoding human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0.8%; Score 13.8; DB 1; Length 17; 88.2%; Pred. No. 9e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 17 BP; 5 A; 8 C; 2 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example 2; Page 179; 418pp; English.
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                                                                                  2001WO-US000666.
2001WO-US000667.
                                                                                                                                                            30-JAN-2001; 2001WO-US000669.
                                2001WO-US000664
2001WO-US000665
                                                                                                                                      2001WO-US000668
                                                                                                                                                                                                                23-MAY-2001; 2001US-00864761
28-AUG-2001; 2001US-0315676P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1397 AGCTGTTGCAGTTTGAG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        17 Agcrerrecadreres
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                         New human kidney tumor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   e.g., liver or bone.
                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2002-479509/51.
                                                                                                                                                                                                                                                                                                  (AEOM-) AEOMICA INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Local Similarity
les 15; Conserv
                                                                                                     30-JAN-2001;
                                                                                  30-JAN-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Homo sapiens
                              30-JAN-2001;
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Query Match

Matches

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Gaps

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2; Indels

amberzyme.

WO200188124-A2.

amberzyme.

ABK18593;

RESULT 1323

22-NOV-2001

DB 1; Length 17;

10017621-381.rng

Page 645

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The invention relates to a nucleic acid molecule (I) which down regulates expression of an Ets-related gene (ERG). (I) is useful for treating conditions selected from cancer, lymphoma, Ewing's sacroma, melanoma, tumour angiogenesis, diabetic retinopathy, macular degeneration, tumour angiogenesis, diabetic retinopathy, macular degeneration, tumour angiogenesis, diabetic retinopathy, macular degeneration, culgaris, angiofibzona of tuberous sclerosis, port-whos stains, Sturge Weber syndrome, Kippel-Trenaunay-Weber syndrome, Osler-Weber-rendu (Weber syndrome, leukaemia, osteoporosis and wound healing. (I) is useful for treating a patient having a condition associated with the level of ERG, by contacting cells of the patient with (I) under conditions suitable for the treatment. Leukaemia or tumour angiogenesis is treated by administering (I) to the patient in conjunction with one or more of other therapies such as radiation or comjunction with one or more of other therapies such as radiation or comjunction with one or more of other therapies such as radiation or comjunction with one or more of other therapies such as radiation or comjunction with a more of other therapies such as radiation or comjunction with a more of other therapies such as radiation or cation such as Mg2+. (I) is useful for reducing ERG activity in a calin, by contacting the expression of ERG, and as diagnostic tool to examine genetic drift and mutations within diseased cells or to detect the presence of ERG RMA in a cell. (I) is useful for specifically craceting genes that share homology with ERG gene or ERG fusion genes. ABK17354-ABK22719 represent nucleic acids, including antisense and carzymatic nucleic acid molecules which regulate expression of ERG, and examine of enzymatic nucleic acides, including antisense and examine the presence of enzymatic nucleic acides, including antisense and enzymatic nucleic acides including antisense and enzymatic nucleic acides including antisense and enzymatic nucleic acides including antisense of ERG Fusion 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0;
                                                                                                                                                                                                                                                                                                                         useful for treating cancer, diabetic retinopathy, macular degeneration,
arthritis, psoriasis, verruca vulgaris and Sturge Weber syndrome.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PAPP-E; human; pregnancy associated plasma protein E; abortive; contraceptive; gene therapy; vaccine; pregnancy; antenatal; diagnosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
                                                                                                                                                                                                                                    Mcswiggen JA, Mclaughlin F, Randi AM;
                                                                                                                                                                                                                                                                                                       Novel polynucleotide which down regulates expression of Ets-related
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         .
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 17 BP; 4 A; 5 C; 6 G; 0 T; 2 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human PAPP-Ea associated 17-mer SEQ ID 576.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   related PCR primers of the invention
                                                                                                                                                                                                                                                                                                                                                                      Claim 4; Page 91; 149pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     17
                                                                                                                  16-MAY-2001; 2001WO-US015866.
                                                                                                                                                   16-MAY-2000; 2000US-00572021.
                                                                                                                                                                                                                                    Von Carlowitz I,
                                                                                                                                                                                   (RIBO-) RIBOZYME PHARM INC. (GLAX ) GLAXO GROUP LTD.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GGAGAUCAGCCUGGACC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      705 GGAGATCAGACTGGAAC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ABS75050 standard; DNA; 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
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                                                                                                                                                                                                                                                                      WPI; 2002-082995/11.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local Similarity
Matches 13; Conserv
                                                 WO200188124-A2.
                 Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            24-DEC-2002
                                                                                  22-NOV-2001
                                                                                                                                                                                                                                    Jarvis T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ABS75050;
                                                                                                                                                                                                                                                                                                                       useful
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          RESULT 1325
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ID ABS
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AC ABS
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DT 24 -:
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DE Hum
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KW PAP
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This invention describes a novel isolated nucleic acid that encodes one of three new isoforms of human pregnancy associated plasma protein E, hpApP-E. The products of the invention have abortive and contraceptive activity and can be used for gene therapy or in a vaccine. The nucleic acid, polypeptide encoded by it, or antibody to the polypeptide can be used in pharmaceutical compositions or vaccines for preventing or aborting pregnancy. PAPP-E is used in the antenatal diagnosis of dysgenetic pregnancies. The nucleic acids are used as probes to assess the level of PAPP-E isoform mRNA in chorionic villus samples, and the antibodies can be used to assess the expression levels of PAPP-E isoform
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               proteins in chorionic villus samples, to diagnose dysgenetic pregnancies antenatally. This sequence represents an oligomer used in scanning the human PAPP-E genes described in the disclosure of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    contraceptive; gene therapy; vaccine; pregnancy; antenatal; diagnosis;
dysgenetic pregnancy; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New isolated nucleic acid encoding an isoform of human pregnancy associated plasma protein E, for preventing or aborting pregnancy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PAPP-E; human; pregnancy associated plasma protein E; abortive;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ö
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 17 BP; 6 A; 2 C; 8 G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 13.8; DB 1
Pred. No. 9e+02;
0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 2; Page 151; 353pp; English
dysgenetic pregnancy; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1011 GAGGGGAGAGCTCAAGC 1027
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  17
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                                                                                                                                                                                                                                                 06-APR-2001; 2001US-00827998.
                                                                                                                                                                                                                                                                                                           26-MAY-2000; 2000US-0207456P.
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88.2%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2002-697817/75.
                                                                                                                                                                                                                                                                                                                                                                                                   (SHAN/) SHANNON M E.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gu Y, Shannon ME;
                                                                                                                     US2002102252-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          JS2002102252-A1
                                                          Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               domo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    24-DEC-2002
                                                                                                                                                                                   01-AUG-2002.
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                                                                                                                                                                                                                                                                                                                                                                         (GUYY/) GU
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ABS75049;
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26-MAY-2000; 2000US-0207456P.

WPI; 2002-684061/74

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human; POSHL 1; SH3 domain; POSH-like signalling protein 1; oncogene;
Rho GTPase; signal transduction; gene expression; cancer; vaccine;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
                                                                                                        New isolated nucleic acid encoding an isoform of human pregnancy associated plasma protein B, for preventing or aborting pregnancy.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human POSHL1 scanning oligonucleotide SEQ ID NO 108.
                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 17 BP; 7 A; 1 C; 8 G; 1 T; 0 U; 0 Other;
                                                                                                                                                       Example 2; Page 150; 353pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1010 AGAGGGGAGAGCTCAAG 1026
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ABV89395 standard; DNA; 17 BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         gene therapy; transgenic; ss.
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2001WO-US000670.
2001US-00864761.
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2001WO-US000665
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(GUYY/) GU Y.
(SHAN/) SHANNON M E.
                                                                          WPI; 2002-697817/75
                                             Shannon ME;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    EP1239051-A2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 RESULT 1327
                                             Gu Y,
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the invention traces to an instance of 730 and acids (S1, ABB8399), a sequence having 65% sequence of 730 and acids (S1, ABB8399), a sequence having 65% sequence identity to (S1), acids (S1, ABB8399), a sequence having 65% sequence identity to (S1), acids (S1, ABB8399), a sequence comprising at least 8 contiguous amino acids. Its appropriate that interacts sequence comprising at least 8 contiguous amino acids. Human POSHL 1 is a proto-oncogene/oncogene product that functions as an adaptor protein that interacts with Rho family small Gypases as well as downstream components of the signal transduction pathway. (I) is useful for identifying a specific binding partner. (I) and nucleic acids (II) for encoding (I) are useful for diagnosing, monitoring disease and treating caused by altered expression of human POSHL1 including diagnosing and treating cancer, they useful in the development of vaccines and (II) is useful in gene therapy. (II) is useful for constructing microarrays which are useful in measuring and for surveying gene expression and creating transgenic non-human animals capable of producing the proteins. The present sequence is that of a scanning oligonucleotide useful in examples of the invention. Note: The present sequence did not form part of the continuous patts of the privent by the European Patent Office
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          .;
0
                                           Novel human SH3 domain (POSH)-like signaling protein 1 polypeptide, POSHL-1, useful for treating disorders associated with decreased expression or
                                                                                                                                                      The invention relates to an isolated SH3 domain (POSH)-like signalling
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human, POSHL 1, SH3 domain, POSH-like signalling protein 1, oncogene; Rho GTPase; signal transduction, gene expression, cancer, vaccine;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
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                                                                                                                Example 2; SEQ ID NO 108; 60pp + Sequence Listing; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 17 BP; 3 A; 2 C; 11 G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     9e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 13.8;
Pred. No. 9e
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2001MO-US000664.
2001MO-US000665.
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2001MO-US000667.
2001MO-US000668.
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2001US-0328205P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Match 0.8%;
Local Similarity 88.2%;
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                                                                         activity of human POSHII.
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30-JAN-2001;
30-JAN-2001;
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30-JAN-2001;
30-JAN-2001;
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2001WO-US000668.
2001WO-US000669.
2001WO-US000670.
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2001WO-US000667.
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30-JAN-2001;
30-JAN-2001;
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       Shannon M;
                                                                                                                                                                                    ABV91270;
                                                                                                                                   Query Match
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ABV91270
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23-MAY-2001; 2001US-00864761. 10-OCT-2001; 2001US-0328205P. (AEOM-) AEOMICA INC Shannon M; g ð The invention relates to an isolated SH3 domain (POSH)-like signalling action [8.1, MBBA399], a sequence having 6fs sequence of 730 amino acids (SI, MBBA399), a sequence having 6fs sequence identity to (SI), (SI) having 9fs deviations, especially conservative substitutions or a fragment of the sequences comprising at least 8 contiguous amino acids. Human POSHL Is a proto-oncogene/oncogene product that functions as an adaptor protein that interacts with Rho family small Grases as well as downstream components of the signal transduction pathway. (I) is useful of for identifying a specific binding partner. (I) and nucleic acids (II) encoding (I) are useful for diagnosing, monitoring disease and treating caused by altered expression of human POSHL1 including diagnosing and caused by altered expression of human POSHL1 including microarrays which useful in gene therapy. (II) is useful for constructing microarrays which are useful for measuring and for surveying gene expression and creating transgenic non-human animals capable of producing the proteins. The present sequence is that of a scanning oligomucleotide useful in examples of the invention. Note: The present sequence did not form part of the prevent by the European Patent Office ö Novel human SH3 domain (POSH)-like signaling protein 1 polypeptide, POSHL-1, useful for treating disorders associated with decreased expression or Gaps Human, POSHL 1, SH3 domain, POSH-like signalling protein 1, oncogene, Rho GTPase; signal transduction, gene expression, cancer, vaccine, ·. DB 1; Length 17; Example 2; SEQ ID NO 280; 60pp + Sequence Listing; English 2; Indels Human POSHL1 scanning oligonucleotide SEQ ID NO 1983. Sequence 17 BP; 2 A; 4 C; 5 G; 6 T; 0 U; 0 Other; 0.8%; Score 13.8; DB 1 88.2%; Pred. No. 9e+02; iive 0; Mismatches 696 GGCACTCAAGGAGATCA 712 ABV91270 standard; DNA; 17 BP 17 GGCACTCCAGAAGATCA 1 gene therapy; transgenic; ss. (first entry) activity of human POSHL1. Best Local Similarity 88.2 Matches 15; Conservative (AEOM-) AEOMICA INC WPI; 2002-684061/74

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The invention relates to an isolated SH3 domain (POSH)-like signalling protein 1 (POSHL 1) polypeptide (I), comprising a sequence of 730 amino acids (S1, ABBB3999), a sequence having 65% sequence identity to (S1), (S1) having 95% deviations, especially conservative substitutions or a fragment of the sequences comprising at least 8 contiguous amino acids. Human POSHL 1 is a prote-oncogene/oncogene product that functions as an adaptor protein that interacts with Rho family small GTPases as well as constream components of the signal transduction pathway. (I) is useful for interitying a specific binding partner. (I) and nucleic acids (II) encoding (I) are useful for diagnosing, monitoring disease and treating caused by altered expression of human POSHL1 including diagnosing and creating caused, they useful in the development of vaccines and (II) is useful in gene therapy. (II) is useful for constructing microarrays which are useful for measuring and for surveying gene expression and creating creating in the development of proteins. The present sequence is that of a scanning oligonucleotide useful in examples of the invention. Note: The present sequence did not form part of the continuant hy, the present sequence information supplied to hermont have a privated the protein or present proteins.
                                                  Novel human SH3 domain (POSH)-like signaling protein 1 polypeptide, POSHL-1, useful for treating disorders associated with decreased expression or activity of human POSHL1.
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                                                                                                                                                                   Example 2; SEQ ID NO 1983; 60pp + Sequence Listing; English.
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88.2%; Pred. No. 9e+02;
lve 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 17 BP; 2 A; 8 C; 6 G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human CLCAl gene enzymatic nucleic acid #808.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          88.2%;
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WPI; 2002-684061/74.
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(RIBO-) RIBOZYME PHARM INC.

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The invention relates to enzymatic nucleic acid molecules that down regulate expression of chloride channel calcium activated 1 (CLCA1) genes cregulate expression of chloride channel calcium activated 1 (CLCA1) genes by cleaving RNA derived from the genes. The nucleic acid sequences are useful as pharmaceutical agents for treating conditions such as chronic obstructive pulmonary disease (CDDD), chronic bronchitis, asthma, cystic fibrosis, obstructive bowel syndrome and any other diseases or conditions consistence, are useful for respond to the levels of CLCA1 in a cell or tissue. The sequences are useful for reducing CLCA1 activity in a cell, chence, are useful for treatment of a patient having a condition acell, consociated with the level of CLCA1, where the invention further comprises treatment, for example, oxygen therapy, bronchodilators, corticosteroids, antibacterials, vaccinations, acetylcysteine and mucokinetic agents. The condition are also used as diagnostic tools to examine genetic drift and mutations within diseased cells or to detect the presence of CLCA1 RNA in a cell. This sequence represents an expresents an expresents and expresents.
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                                                                                                                                                        Enzymatic polynucleotide that down regulates expression of chloride channel calcium activated gene, useful for treating Chronic obstructive pulmonary disease (COPD), chronic bronchitis and asthma.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
                                                           Szymkowski DE;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 17 BP; 6 A; 5 C; 4 G; 0 T; 2 U; 0 Other;
                                                             Mckenzie T, Ayers D,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human CLCA1 gene enzymatic nucleic acid #1498.
                                                                                                                                                                                                                                               Claim 4; Page 70; 152pp; English.
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                                                             Mcswiggen J,
(SYNT ) SYNTEX USA LLC. (THOM/) THOMPSON J.
                                                                                                                        WPI; 2002-217145/27.
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                                                                 Thompson J,
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                                                                                      Grupe A;
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The invention relates to enzymatic nucleic acid molecules that down regulate expression of chloride channel calcium activated 1 (CLCA1) genes or regulate expression of chloride channel calcium activated 1 (CLCA1) genes by cleaving RNA derived from the genes. The nucleic acid sequences are useful as pharmaceutical agents for treating conditions such as chronic cobstructive pulmonary disease (CDCD), chronic bronchitis, asthma, cystic fibrosis, obstructive bowel syndrome and any other diseases or conditions tissue. The sequences are useful for reducing CLCA1 activity in a cell, chence, are useful for treatment of a patient having a condition associated with the level of CLCA1, where the invention further comprises the use of one or more therapies under conditions suitable for the treatment, for example, oxygen therapy, bronchodilators, corticosteroids, antibacterials, vaccinations, acetylcysteine and mucokinetic agents. The nucleic acids of the invention are also used as diagnostic tools to camine genetic drift and mutations within diseased cells or to detect the presence of CLCA1 RNA in a cell. This sequence represents an
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human; chloride channel calcium activated 1; CLCA1; ss; antiasthmatic; antinflammatory; chronic obstructive pulmonary disease; COPD; asthma; chronic bronchitis; cystic fibrosis; obstructive bowel syndrome; oxygen therapy; bronchodilator; corticosteroid; vaccination; mucokinetic;
                                                                                      Enzymatic polynucleotide that down regulates expression of chloride channel calcium activated gene, useful for treating Chronic obstructive pulmonary disease (COPD), chronic bronchitis and asthma.
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Szymkowski DE;
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Grupe A;
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 Mcswiggen J, Mckenzie T, Ayers D,
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                                                                                                                                                                        Claim 4; Page 96; 152pp; English.
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     Thompson J,
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                         Grupe A;
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WPI; 2002-217145/27

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Sequence 17 BP; 4 A; 5 C; 4 G; 0 T; 4 U; 0 Other;
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                                                                                                                                                                            Claim 4; Page 70; 152pp; English
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nes 12; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          West Nile Virus.
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(MCSW/) N
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        RESULT 1333
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HANNER KARAKAR KARAKAR
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AAGCAGGCCAGCUUUUC 17

BP.

(first entry)

Mcswiggen JA;

Blatt L,

WPI; 2002-706994/76.

MCSWIGGEN J A.

0.8%; Score 13.8; DB 1; Length 17; 70.6%; Pred. No. 9e+02; ve 3; Mismatches 2; Indels

70.6%;

Conservative

The invention relates to nucleic acid molecules that modulate replication of the West Nile Virus (WNV). The nucleic acid molecules are useful for treating a condition related to WNV infection e.g. pancreatitis, invection e.g. pancreatitis, liver failure, hepatocalfular carcinoma or cirrhosis. The nucleic acid molecule is selected from the group of ribozymes consisting of Hammerhead, Inozyme, G-cleaver, DNAzyme, Amberzyme and Zinzyme. The nucleic acid molecules further comprise at least five ribose residues, at least ten 2'-O-methyl modifications, phosphorothioate linkages on at least three of the 5' terminal nucleotides and a 3' end modification of a 3'-3' inverted abasic moiety. Nucleic acid molecules SEQ ID NO 1 to 37080 are claimed; however, SEQ ID NO 2194-2206 and 17502-17514 are not given in the specification. The present sequence is that of a nucleic acid The invention relates to nucleic acid molecules that modulate replication of the West Nile Virus (WNV). The nucleic acid molecules are useful for treating a condition related to WNV infection e.g. pancreatitis, New nucleic acid molecule that modulates replication of West Nile Virus (WNV), useful for treating a condition related to WNV infection e.g. pancreatitis, meningitis, hepatocellular carcinoma or cirrhosis. Gaps WNV; West Nile Virus; antiinflammatory; cytostatic; hepatotropic; virucide; neuroprotective; antibacterial; replication; pancreatitis; encephalitis; myocarditis; meningitis; infection; hepatitis; liver failure; cancer; cirrhosis; Hammerhead; Inozyme; DNAzyme; pancreatitis, meningitis, hepatocellular carcinoma or cirrhosis. . 0 Score 13.8; DB 1; Length 17; Pred. No. 9e+02; 0; Mismatches 2; Indels Indels WNV minus strand Inozyme substrate SEQ ID NO 10400. Sequence 17 BP; 1 A; 2 C; 9 G; 0 T; 5 U; 0 Other; Claim 23; SEQ ID NO 10400; 495pp; English. Claim 23; SEQ ID NO 6745; 495pp; English ·. 447 ACN10397 standard; RNA; 17 BP. 19-OCT-2001; 2001WO-US048350. 20-OCT-2000; 2000US-0242411P. 17 ACCAACCCCCACGCATG 1 0.8%; 88.2%; 431 ACCATCCCCCACGCAAG RIBOZYME PHARM INC. (first entry) molecule of the invention Query Match 0.8 Best Local Similarity 88.2 Matches 15; Conservative Blatt L, Mcswiggen JA; Amberzyme; Zinzyme; ss (BLAT/) BLATT L, (MCSW/) MCSWIGGEN J A. WPI; 2002-706994/76. West Nile Virus. WO200268637-A2. 22-APR-2004 06-SEP-2002. ACN10397; (RIBO-) RESULT 1334 ACN10397 à g The invention relates to enzymatic nucleic acid molecules that down regulate expression of chloride channel calcium activated 1 (CLCA1) genes by cleaving RNA derived from the genes. The nucleic acid sequences are useful as pharmaceutical agents for treating conditions such as chronic cobstructive pulmonary disease (CDD), chronic bronchitis, asthma, cystic fibrosis, obstructive bowel syndrome and any other diseases or conditions that are related to or will respond to the levels of CLCA1 in a cell or tissue. The sequences are useful for reducing CLCA1 activity in a cell, hence, are useful for treatment of a patient having a condition associated with the level of CLCA1, where the invention further comprises the use of one or more therapies under conditions suitable for the treatment, for example, oxygen therapy, bronchodilators, corticosteroids, antibacterials, vaccinations, acetylcysteine and mucokinetic agents. The nucleic acids of the invention are also used as diagnostic tools to examine genetic drift and mutations within diseased cells or to detect the presence of CLCA1 RNA in a cell. This sequence represents an ·. Enzymatic polynucleotide that down regulates expression of chloride channel calcium activated gene, useful for treating Chronic obstructive pulmonary disease (COPD), chronic bronchitis and asthma. New nucleic acid molecule that modulates replication of West Nile Virus (MNV), useful for treating a condition related to MNV infection e.g. Gaps WNV; West Nile Virus; antiinflammatory; cytostatic; hepatotropic; virucide; neuroprotective; antibacterial; replication; pancreatitis; encephalitis; myocarditis; meningitis; infection; hepatitis; liver failure; cancer; cirrhosis; Hammerhead; Inozyme; DNAzyme; 0;

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encephalitis, myocarditis, meningitis, neurologic infection, hepatitis, liver failure, hepatocellular carcinoma or cirrhosis. The nucleic acid molecule is selected from the group of ribozymes consisting of Hammerhead, Inozyme, G-cleaver, DNAzyme, Amberzyme and Zinzyme. The least ten 2·O-mehbyl modifications, phosphorothicate linkages on at least three of the 5' terminal nuclectides and a 3' end modification of 3'-3' inverted abasic moiety. Nucleic acid molecules SEQ ID NO 1 to 37080 are claimed, however, SEQ ID NO 2194-2206 and 17502-17514 are not given molecule of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New nucleic acid molecule that modulates replication of West Nile Virus (WNV), useful for treating a condition related to WNV infection e.g. pancreatitis, meningitis, hepatocellular carcinoma or cirrhosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WNV; West Nile Virus; antiinflammatory; cytostatic; hepatotropic; virucide; neuroprotective; antibacterial; replication; pancreatitis; encephalitis; myocarditis; meningitis; infection; hepatitis; liver failure; cancer; cirrhosis; Hammerhead; Inozyme; DNAzyme;
                                                                                                                                                                                                                                          Gaps
                                                                                                                                                                                                                                          ;
0
                                                                                                                                                                                                            0.8%; Score 13.8; DB 1; Length 17; 12.4%; Pred. No. 9e+02; ve 1; Mismatches 2; Indels
                                                                                                                                                                               G; 0 T; 2 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WNV Amberzyme substrate SEQ ID NO 5561.
                                                                                                                                                                                                                                                                         433 CATCCCCCACGCAAGAT 449
                                                                                                                                                                                                                                                                                                     CAACCCCCACGCAUGAU 17
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                                                                                                                                                                                   Sequence 17 BP; 5 A; 8 C; 2
                                                                                                                                                                                                                             82.4%;
                                                                                                                                                                                                                                                                                                                                                                              ACN05558 standard; RNA; 17
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                                                                                                                                                                                                                                            14; Conservative
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(MCSW/) MCSWIGGEN J A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Amberzyme; Zinzyme; ss
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                                                                                                                                                                                                                                Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    West Nile Virus
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                                                                                                                                                                                                                  Query Match
                                                                                                                                                                                                                                                                                                                                              RESULT 1335
ACN05558/c
                                                                                                                                                                                                                                             Matches
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treating a condition related to WNV infection e.g. parcreatists.

C encephalitis, myocarditis, meningitis, neurologic infection, hepatitis,

C inver failure, hepatocellular carcinoma or cirrhosis. The nucleic acid

C liver failure, hepatocellular carcinoma or cirrhosis. The nucleic acid

C Hammerhead, Inozyme, G-cleaver, DNAzyme, Amberzyme and Zinzyme. The

CC Hammerhead, Inozyme, G-cleaver, DNAzyme, Amberzyme and Zinzyme, The

CC least three of the 5' terminal nucleotides and a 3' end modification of a collect three of the 5' terminal nucleotides and a 3' end modification of a 3'-3' inverted abasic moiecy. Nucleic acid molecules SEQ ID NO 1 to 37080

CC are claimed; however, SEQ ID NO 2194-2206 and 17502-17514 are not given in the specification. The present sequence is that of a nucleic acid

X molecule of the invention
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3'-3' inverted abasic moiety. Nucleic acid molecules SEQ ID NO 1 to 37080 are claimed, however, SEQ ID NO 2194-2206 and 17502-17514 are not given in the specification. The present sequence is that of a nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New nucleic acid molecule that modulates replication of West Nile Virus (WNV), useful for treating a condition related to WNV infection e.g. pancreatitis, meningitis, hepatocellular carcinoma or cirrhosis.
                                                                                                                                                                    Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WNV; West Nile Virus; antiinflammatory; cytostatic; hepatotropic; virucide; neuroprotective; antibacterial; replication; pancreatitis; encephalitis; myocarditis; meningitis; infection; hepatitis; liver failure; cancer; cirrhosis; Hammerhead; Inozyme; DNAzyme;
                                                                                                                                                                    ..
                                                                                                                              0.8%; Score 13.8; DB 1; Length 17; 18.2%; Pred. No. 9e+02;
                                                                                                                                                                    Indels
                                                                                                                                                                  5,
                                                                                             Sequence 17 BP; 3 A; 2 C; 6 G; 0 T; 6 U; 0 Other;
                                                                                                                                                                      0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 claim 23; SEQ ID NO 3583; 495pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                         Zinzyme substrate SEQ ID NO 3583.
                                                                                                                                                                                                       477 ATCACTACCAGCTGACA 493
                                                                                                                                                                                                                                                                                                                                  BP.
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                                                                                                                                                     88.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (RIBO-) RIBOZYME PHARM INC.
                                                                                                                                                                                                                                            17 ATCATTACCAGCCGACA
                                                                                                                                                                                                                                                                                                                                    ACN03580 standard; RNA; 17
                                                                                                                                                                                                                                                                                                                                                                                                         (first entry)
                                                               molecule of the invention
                                                                                                                                    Query Match 0.8
Best Local Similarity 88.2
Matches 15, Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Blatt L, Mcswiggen JA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       West Nile Virus.
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Sequence 17 BP; 4 A; 1 C; 8 G; 0 T; 4 U; 0 Other;

The invention relates to nucleic acid molecules that modulate replication of the West Nile Virus (WNV). The nucleic acid molecules are useful for treating a condition related to WNV infection e.g. pancreatitis, encephalitis, wyocarditis, meningitis, neurologic infection, hepatitis, liver failure, hepatocellular carcinoma or cirrhosis. The nucleic acid molecule is selected from the group of ribozymes consisting of Hammerhead, Inozyme, G-cleaver, DNAzyme, Amberzyme, and Zinzyme. The nucleic acid molecules further comprise at least five ribose residues, at least ten 2'-O-methyl modifications, phosphorothioate linkages on at least three of the 5' terminal nucleotides and a 3' end modification of a

claim 23; SEQ ID NO 5561; 495pp; English.

WNV; West Nile Virus; antiinflammatory; cytostatic; hepatotropic; virucide; neuroprotective; antibacterial; replication; pancreatitis; encephalitis; myocarditis; meningitis; infection; hepatitis; liver fallure; cancer; cirrhosis; Hammerhead; Inozyme; DNAzyme;

WNV minus strand DNAzyme substrate SEQ ID NO 13858.

(first entry)

22-APR-2004

ACN13855;

ACN13855 standard; RNA; 17 BP

RESULT 1338

ACN13855/

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New nucleic acid molecule that modulates replication of West Nile Virus
                                         Gaps
                                                                                                                                                                                                                                                                                                                                           WNV; West Nile Virus; antiinflammatory; cytostatic; hepatotropic; virucide; neuroprotective; antibacterial; replication; pancreatitis; encephalitis; myocarditis; meningitis; infection; hepatitis; liver failure; cancer; cirrhosis; Hammerhead; Inozyme; DNAzyme;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (WNV), useful for treating a condition related to WNV infection e.g. pancreatitis, meningitis, hepatocellular carcinoma or cirrhosis.
                                         .
     Length 17;
             ,
9e+02;
----- 2; Indels
                                                                                                                                                                                                                                                                                                           WNV minus strand Inozyme substrate SEQ ID NO 10761.
0.8%; Score 13.8; DB 1;
88.2%; Pred. No. 9e+02;
.ve 0; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 17 BP; 2 A; 2 C; 6 G; 0 T; 7 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 23; SEQ ID NO 10761; 495pp; English.
                                                                       1491 TCCTGACACTACTTCCA 1507
                                                                                                                                                                                                    ACN10758 standard; RNA; 17 BP
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                                                                                                         17 recadacacierrica 1
                  88.2%;
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                                                                                                                                                                                                                                                                        22-APR-2004 (first entry)
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                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mcswiggen JA;
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                    Similarity
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                                       15;
Query Match
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                      Best Local
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                                   Matches
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ACN10758/c
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of the West Nile virus (WWV). The mucleic acid molecules are useful for treating a condition related to WNV infection e.g. pancreatitis, encephalitis, myocarditis, meningitis, neurologic infection, hepatitis, liver failure, hepatocellular carcinoma or cirrhosis. The mucleic acid molecule is selected from the group of ribozymes consisting of Hammerhead, Inozyme, G-Cleaver, DNAzyme, Amberzyme and Zinzyme. The nucleic acid molecules further comprise at least five ribose residues, at least three of the 5' terminal nucleotides and a 3' end modification of a 3'-3' inverted abasic moiety. Nucleic acid molecules SEQ ID NO 110 37080 are claimed, however, SEQ ID NO 2194-2206 and 17502-17514 are not given in the specification. The present sequence is that of a nucleic acid
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88.2%; Pred. No. 9e+02;
ve 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 17 BP; 2 A; 3 C; 5 G; 0 T; 7 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 23; SEQ ID NO 13858; 495pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ACN12120 standard; RNA; 17 BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      nolecule of the invention
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                                                                                                                                                                               Amberzyme; Zinzyme; ss
                                                                                                                                                                                                                                                                                                                                                                                             (BLAT/) BLATT L.
(MCSW/) MCSWIGGEN J A.
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les 15; Conserv
                                                                                                                                                                                                            West Nile Virus.
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ACN12120;

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Gaps

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Indels

5;

0.8%; Score 13.8; DB 1 88.2%; Pred. No. 9e+02; ative 0; Mismatches

CATCAACATGCACAACG 910

Conservative

15;

Matches

Similarity

Query Match Best Local &

17 CATCAACATGCCAAACG 1

g ò

DB 1; Length 17;

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treating a condition related to WNV infection e.g. pancreatitis, encephalitis, myocarditis, meningitis, encephalitis, myocarditis, encephalitis, phosphorothioate linkages on at least three of the 5' terminal nucleotides and a 3' end modification of a 3' inverted abasic modety. Nucleic acid molecules SEQ ID NO 194-2206 and 17502-17514 are not given in the specification. The present sequence is that of a nucleic acid molecule of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The invention relates to nucleic acid molecules that modulate replication
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New nucleic acid molecule that modulates replication of West Nile Virus (WNV), useful for treating a condition related to WNV infection e.g. pancreatitis, meningitis, hepatocellular carcinoma or cirrhosis.
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encephalitis; myocarditis; meningitis; infection; hepatitis;
liver failure; cancer; cirrhosis; Hammerhead; Inozyme; DNAzyme;
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0
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2%; Pred. No. 9e+02;
0; Mismatches 2; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 17 BP; 5 A; 9 C; 2 G; 0 T; 1 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 23; SEQ ID NO 10398; 495pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      431 ACCATCCCCCACGCAAG 447
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                                                                                                                                                                                                                                            19-OCT-2001; 2001WO-US048350.
                                                                                                                                                                                                                                                                                            20-OCT-2000; 2000US-0242411P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ACN10396 standard; RNA; 17
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                                                                                                                                                                                                                                                                                                                                                                                                                                        Blatt L, Mcswiggen JA;
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                                                                                                                                                                                                                                                                                                                                         (RIBO-) RIBOZYME PHARM
                                                    Amberzyme; Zinzyme; ss
                                                                                                                                                                                                                                                                                                                                                               (BLAT/) BLATT L.
(MCSW/) MCSWIGGEN J A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2002-706994/76.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity
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                                                                                                 West Nile Virus.
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                                                                                                                                                   WO200268637-A2.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ACN10396
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New nucleic acid molecule that modulates replication of West Nile Virus (WNV), useful for treating a condition related to WNV infection e.g. pancreatitis, meningitis, hepatocellular carcinoma or cirrhosis.
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                                                                                                                          WNV; West Nile Virus; antiinflammatory; cytostatic; hepatotropic; virucide; neuroprotective; antibacterial; replication; pancreatitis; encephalitis; myocarditis; meningitis; infection; hepatitis; liver failure; cancer; cirrhosis; Hammerhead; Inozyme; DNAzyme;
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                                                                               WNV minus strand Inozyme substrate SEQ ID NO 12123.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0.8%; Score 13.8; DB 176.5%; Pred. No. 9e+02; Ive 2; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 23; SEQ ID NO 12123; 495pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                477 ATCACTACCAGCTGACA 493
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1 Auchunaccascosaca 17
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                                                                                                                                                                                                                                                                                                                                                                                                                      19-OCT-2001; 2001WO-US048350
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RIBOZYME PHARM INC
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                                  (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Local Similarity 76.5
nes 13; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mcswiggen JA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           BLATT L.
MCSWIGGEN J A.
                                                                                                                                                                                                                              Amberzyme; Zinzyme; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2002-706994/76.
                                                                                                                                                                                                                                                                        West Nile Virus.
                                                                                                                                                                                                                                                                                                                          WO200268637-A2.
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                                  22-APR-2004
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(MCSW/)
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WPI; 2002-706994/76
                                                                                 Best Local Similarity
                                                                                                                                   West Nile Virus
                                                                                                                                       WO200268637-A2.
  06-SEP-2002.
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                                                                                                          ACN10409;
                                                                               Query Match
                                                                                                 RESULT 1342
ACN10409/c
                                                                                   Matches
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The invention relates to nucleic acid molecules that modulate replication of the West Nile Virus (WNV). The nucleic acid molecules are useful for treating a condition related to WNV infection e.g. pancreatitis, morphalitis, mycoarditis, meningitis, neurologic infection, hepatitis, liver failure, hepatocellular carcinoma or cirrhosis. The nucleic acid molecule is selected from the group of ribozymes consisting of Hammerhead, Inozyme, G-cleaver, DNAzyme, Amberzyme and Zinzyme. The nucleic acid molecules further comprise at least five ribose residues, at least ten 2'-O-methyl modifications, phosphorothioate linkages on at least three of the 5' terminal nucleotides and a 3' end modification of a 3'-3' inverted abasic moiety. Nucleic acid molecules SEQ ID NO 1 to 37080 are claimed; however, SEQ ID No 2194-2206 and 17502-17514 are not given in the specification. The present sequence is that of a nucleic acid
                                                                                                                                                                                           New nucleic acid molecule that modulates replication of West Nile Virus (WNV), useful for treating a condition related to WNV infection e.g. pancreatitis, meningitis, hepatocellular carcinoma or cirrhosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WNV; West Nile Virus; antiinflammatory; cytostatic; hepatotropic; viruoide; neuroprotective; antibacterial; replication; pancreatitis; encephalitis; mycoarditis; meningitis; infection; hepatitis; liver failure; cancer; cirrhosis; Hammerhead; Inozyme; DNAzyme;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0.8%; Score 13.8; DB 1; Length 17; 88.2%; Pred. No. 9e+02; ive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 17 BP; 4 A; 8 C; 1 G; 0 T; 4 U; 0 Other;
                                                                                                                                                                                                                                                                                           Claim 23; SEQ ID NO 10412; 495pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WNV Inozyme substrate SEQ ID NO 1663.
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  RIBOZYME PHARM INC.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Local Similarity 88.2
les 15; Conservative
                                                                                                  Mcswiggen JA;
                                                (MCSW/) MCSWIGGEN J A.
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                                                                                                                                                WPI; 2002-706994/76
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                                                                                                Blatt L,
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(RIBO-) H
(BLAT/) H
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             of the West Nile Virus (WNV). The nucleic acid molecules are useful for treating a condition related to WNV infection e.g. pancreatins, encephalitis, myocarditis, meningitis, neurologic infection, hepatitis, liver failure, hepatocellular carcinoma or cirrhosis. The nucleic acid molecule is selected from the group of ribozymes consisting of Hammerhead, Inoxyme, G-cleaver, DNAzyme, Amberzyme, and Zinzyme. The nucleic acid molecules further comprise at least five ribose residues, at least the Z-O-methyl modifications, phosphorothicate linkages on at least three of the 5' terminal nucleotides and a 3' end modification of a 3'-3' inverted abasic moiety. Nucleic acid molecules SEQ ID NO 1 to 37080 in the specification. The present sequence is that of a nucleic acid nolecule of the invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The invention relates to nucleic acid molecules that modulate replication
                                                                                                                                                                                                                                                                                                                                                             nucleic acid molecule that modulates replication of West Nile Virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
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                                                                                                                                                                                                                                                                                                                                                                                 (WNV), useful for treating a condition related to WNV infection e.g. pancreatitis, meningitis, hepatocellular carcinoma or cirrhosis.
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88.2%; Pred. No. 9e+02;
7ative 0; Mismatches 2; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 23; SEQ ID NO 10399; 495pp; English.
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ACN10409 standard; RNA; 17 BP
                                                                 19-OCT-2001; 2001WO-US048350
                                                                                                               20-OCT-2000; 2000US-0242411P
                                                                                                                                                             (RIBO-) RIBOZYME PHARM INC.
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                                                                                                                                                                                                                                                               Blatt L, Mcswiggen JA;
                                                                                                                                                                                                          (MCSW/) MCSWIGGEN J A.
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(RIBO-) RIBOZYME PHARM INC.

(BLAT/) BLATT L. (MCSW/) MCSWIGGEN J A.

19-OCT-2001; 2001WO-US048350 20-OCT-2000; 2000US-0242411P

Blatt L, Mcswiggen JA;

WPI; 2002-706994/76.

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New nucleic acid molecule that modulates replication of West Nile Virus (WNV), useful for treating a condition related to WNV infection e.g. pancreatitis, meningitis, hepatocellular carcinoma or cirrhosis.
                                                                                                                                                                           Sequence 17 BP; 5 A; 6 C; 2 G; 0 T; 4 U; 0 Other;
                                       SEQ ID NO 1663; 495pp; English
                                                                                                                                                                                                                                                                                                                                  substrate SEQ ID NO 6744.
                                                                                                                                                                                                                           1136 ACTACTCCACTCAGATT 1152
                                                                                                                                                                                                                                      1 ACUACUCCACACAGGUU 17
                                                                                                                                                                                                                                                                                    BP.
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                                                                                                                                                                                                     64.78;
                                                                                                                                                                                                                                                                                   ACNO6741 standard; RNA; 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (RIBO-) RIBOZYME PHARM INC.
                                                                                                                                                                                                                                                                                                                   (first entry)
                                                                                                                                                                                                           Matches 11; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (MCSW/) MCSWIGGEN J A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2002-706994/76.
                                                                                                                                                                                                    Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                  West Nile Virus
                                                                                                                                                                                                                                                                                                                                                                                                                  WO200268637-A2.
                                                                                                                                                                                                                                                                                                                                  WNV Amberzyme
                                                                                                                                                                                                                                                                                                                   22-APR-2004
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                                       Claim 23;
                                                                                                                                                                                                                                                                                                    ACN06741;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Blatt L,
                                                                                                                                                                                            Query Match
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The invention relates to nucleic acid molecules that modulate replication of the West Nile Virus (WNV). The nucleic acid molecules are useful for treating a condition related to WNV infection e.g. pancreatis.

Inver failure, hepatocellular carcinoma or cirrhosis. The nucleic acid molecule is selected from the group of ribozymes consisting of Hammerhead, Inozyme, G-cleaver, DNAzyme, Amberzyme and Zinzyme. The nucleic acid molecules further comprise at least five ribose residues, at least ten 2'-O-methyl modifications, phosphorothicate linkages on at least three of the 5' terminal nuclectides and a 3' end modification of a 3'-3' inverted abasic moiety. Nucleic acid molecules SEQ ID NO 1 to 37080 are claimed; however, SEQ ID NO 2194-2206 and 17502-17514 are not given in the specification. The present sequence is that of a nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New nucleic acid molecule that modulates replication of West Nile Virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WNV, West Nile Virus; antiinflammatory; cytostatic; hepatotropic; virucide; neuroprotective; antibacterial; replication; panoreatitis; encephalitis; myocarditis; meningitis; infection; hepatotis; liver failure; cancer; cirrhosis; Hammerhead; Inozyme; DNAzyme;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (WNV), useful for treating a condition related to WNV infection e.g. pancreatitis, meningitis, hepatocellular carcinoma or cirrhosis.
                                                                                                                                                                                                                                                                                                                                                                                                     Score 13.8; DB 1; Length 17; Pred. No. 9e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                      2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WNV minus strand Zinzyme substrate SEQ ID NO 13438.
                                                                                                                                                                                                                                                                                                                                                            Sequence 17 BP; 1 A; 2 C; 9 G; 0 T; 5 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                    0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  claim 23; SEQ ID NO 13438; 495pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  432 CCATCCCCCACGCAAGA 448
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ACN13435 standard; RNA; 17 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    20-OCT-2000; 2000US-0242411P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         19-OCT-2001; 2001WO-US048350.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Н
                                                                                                                                                                                                                                                                                                                                                                                                                                88.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (RIBO-) RIBOZYME PHARM INC. (BLAT/) BLATT L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  17 CCAACCCCCACGCATGA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
                                                                                                                                                                                                                                                                                                                   molecule of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                        15, Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Blatt L, Mcswiggen JA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Amberzyme; Zinzyme; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (MCSW/) MCSWIGGEN J A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2002-706994/76.
                                                                                                                                                                                                                                                                                                                                                                                                                                Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 West Nile Virus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WO200268637-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                22-APR-2004
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ACN13435;
                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RESULT 1345
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ACN13435,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              g
                5
                                                                                                                                                                       The invention relates to nucleic acid molecules that modulate replication of the West Nile Virus (NNV). The nucleic acid molecules are useful for treating a condition related to WNV infection e.g. pancreatitis, mocarditis, muscipalitis, muscipalitis, muscipalitis, muscipalitis, muscarditis, mentagitis, neurologic infection, hepatitis, liver failure, hepatocellular carcinoma or cirrhosis. The nucleic acid molecule is selected from the group of ribozymes consisting of Hammerhead, Inozyme, G-cleaver, DNAzyme, Amberzyme and Zinzyme. The nucleic acid molecules further comprise at least five ribose residues, at least three of the 5' terminal nucleotides and a 3' end modification of a 3'-3' inverted abasic moiety. Nucleic acid molecules SEQ ID NO 1 to 37080 are claimed; however, SEQ ID NO 2194-2206 and 17502-17514 are not given molecule of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ö
                                    New nucleic acid molecule that modulates replication of West Nile Virus (WNV), useful for treating a condition related to WNV infection e.g. pancreatitis, meningitis, hepatocellular carcinoma or cirrhosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WNV; West Nile Virus; antiinflammatory; cytostatic; hepatotropic; virucide; neuroprotective; antibacterial; replication; panoreatitis; encephalitis; myocarditis; meningitis; infection; hepatitis; liver failure; cancer; cirrhosis; Hammerhead; Inozyme; DNAzyme;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             .
0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0.8%; Score 13.8; DB 1; Length 17; 4.7%; Pred. No. 9e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               4; Mismatches
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The invention relates to mucleic acid molecules that modulate replication of the West Nile Virus (WNV). The nucleic acid molecules are useful for treating a condition related to WNV infection e.g. pancreatitis, mencephalitis, meniopiis, meniopiis, neurologic infection, hepatitis, liver failure, hepatocellular carcinoma or cirrhosis. The nucleic acid molecule is selected from the group of ribozymes consisting of Hammerhead, Inozyme, G-cleaver, DNAzyme, Amberzyme and Zinzyme. The

Claim 23; SEQ ID NO 6744; 495pp; English

. 0

Gaps

0

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proteins and their coding sequences: MD24, MD27, MD212. MD23 is encoded at chromosome 7q22.1, MD24 is encoded at chromosome 6p21.3-22.2, MD24, MD21 is encoded at chromosome 6p21.3-22.2, MD27 is encoded at chromosome 16p1.2 and MD212 sequences are useful in therapy, or in manufacturing a medicament for treating or preventing a disorder or in manufacturing a medicament for treating or preventing a disorder. MD24, MD27, e.g. cancer or developmental disorders. The nucleic also and proteins are also useful for diagnoshing or monitoring a disease caused by altered expression of MD23, MD24, MD27, or MD212. The nucleic acids and proteins are also useful for diagnoshing or monitoring a disease caused by altered expression of MD23, MD24, MD27, or MD212. The nucleic acids can also be used as probes to detect and characterize gross acids can also be used as probes to detect and characterize gross alterations in MD23, MD24, MD27, or MD212 genetic locus. The probes are useful in constructing microarrays for measuring gene expression. The proteins are useful as therapeutic agents for gene therapy or as
least ten 2'-O-methyl modifications, phosphorothicate linkages on at least three of the 5' terminal nucleotides and a 3' end modification of a 3'-3' inverted abasic molety. Nucleic acid molecules SED ID NO 1 to 37080 are claimed; however, SEQ ID NO 2194-2206 and 17502-17514 are not given in the specification. The present sequence is that of a nucleic acid molecule of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Cytostatic; immunostimulant; gene therapy; vaccine; human;
zinc finger protein; MDZ3; MDZ4; MDZ7; MDZ1; chromosome 7q22.1;
chromosome 6p21.3-22.2; chromosome 16p11.2; chromosome 15q26.1; cancer;
developmental disorder; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New zinc finger-containing proteins and nucleic acids, useful in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MDZ3,
                                                                                                                                                                                                                                                             Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The present invention relates to novel human zinc finger-containing
                                                                                                                                                                                                                                                           ;
                                                                                                                                                                                                                 DB 1; Length 17;
                                                                                                                                                                                                                                                         2; Indels
                                                                                                                                                                            Sequence 17 BP; 4 A; 3 C; 6 G; 0 T; 4 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human MDZ7 scanning oligonucleotide SEQ ID 4421.
                                                                                                                                                                                                               Score 13.8, DB 1
Pred. No. 9e+02;
                                                                                                                                                                                                                                                           0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 8; SEQ ID NO 4421; 103pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MDZ4, MDZ7 or MDZ12, e.g. cancer
                                                                                                                                                                                                                                                                                              1137 CTACTCCACTCAGATTG 1153
                                                                                                                                                                                                                                                                                                                                                                                                                                     ADB03435 standard; DNA; 17 BP
                                                                                                                                                                                                                   0.8%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gu Y, Nguyen C;
                                                                                                                                                                                                                                                                                                                                      CTACTCCACACAGGTTG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  20-NOV-2003 (first entry)
                                                                                                                                                                                                                                                         15; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2003-423107/40.
                                                                                                                                                                                                                                      Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Homo sapiens.
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                                                                                                                                                                                                                                                                                                                                    17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           acid molecule or an enzymatic nucleic acid molecule, that modulates expression of a nucleic acid molecule encoding HER2, K-Ras, H-Ras, N-Ras, unman immundeficioncy virus (HIV) or a component of HIV. The nucleic acid molecule of the invention has cytostatic, anti-HIV, and anti-rheumatic activity. The nucleic acid molecules are useful for reducing also useful for reducing breast, ovarian, colorectal, lung, prostate, bladder, or pancreatic cancer, and HIV infection, and AIDS. The sequences shown in ABZ59889 - ABZ62216, ABZ64544 - ABZ65531, ABZ66520 - ABZ66524, ABZ66530 - ABZ66585 represent substrate/target sequences for the human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The invention relates to a novel short interfering RNA (siRNA) nucleic
                                                                 Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Novel short interfering RNA and enzymatic nucleic acid useful for treating cancer, modulates the expression of a nucleic acid encoding HBR2, K-Ras, H-Ras, N-Ras, and human deficiency virus sequences.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
                                                                                                                                                                                                                                                                                                                                 Human; ribozyme; short interfering RNA; siRNA; HER2; K-Ras;
enzymatic nucleic acid; H-Ras; N-Ras; HIV; cytostatic; anti-HIV;
anti-rheumatic; cancer; AIDS; ss.
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0
                             Score 13.8; DB 1; Length 17;
Pred. No. 9e+02;
0; Mismatches 2; Indels
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18.2%; Pred. No. 9e+02;
ve 0; Mismatches 2; Indels
Sequence 17 BP; 0 A; 10 C; 3 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 17 BP; 2 A; 4 C; 9 G; 0 T; 2 U; 0 Other;
                                                                                                                                                                                                                                                                                                   Human K-Ras DNAzyme substrate #17.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 58; Page 85; 185pp; English.
                                                              ·.
                                                                                              937
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                                                                                                                                                                                                         BP.
                                                                                                                          ccrerrcccecreccc 17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        29-MAY-2001; 2001US-0294140P.
06-JUN-2001; 2001US-0296249P.
10-SEP-2001; 2001US-0318471P.
                               0.8%;
                                              88.28;
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                                                                                            921 CCTGTTCCAGCTGCTCC
                                                                                                                                                                                                       ABZ59905 standard; RNA; 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (RIBO-) RIBOZYME PHARM INC.
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                                                                                                                                                                                                                                                                    (first entry)
                                                              15; Conservative
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                                               Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                               WO200297114-A2
                                                                                                                                                                                                                                                                                                                                                                                                 Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mcswiggen J;
                                                                                                                                                                                                                                                                      21-MAR-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                             05-DEC-2002
                                                                                                                                                                                                                                     ABZ59905;
                               Query Match
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The invention relates to a novel short interfering RNA (siRNA) nucleic acid molecule or an enzymatic nucleic acid molecule, that modulates expression of a nucleic acid molecule encoding HER2, K.Ras, H.Ras, N.Ras, human immunodeficiency virus (HIV) or a component of HIV. The nucleic acid molecule of the invention has cytostatic, anti-HIV, and anti-rheumatic activity. The nucleic acid molecules are useful for reducing HER2, K-Ras, H-Ras, and HIV activity in a cell. The nucleic acids are also useful for treating breast, ovarian, colorectal, lung, prostate, bladder, or pancreatic cancer, and HIV infection, and AIDS. The sequences shown in ABZ59889 - ABZ62216, ABZ65531, ABZ65520 - ABZ65524, ABZ65530 - ABZ65585 represent substrate/target sequences for the human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  treating cancer, modulates the expression of a nucleic acid encoding HER2, K-Ras, H-Ras, N-Ras, and human deficiency virus sequences.
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Novel short interfering RNA and enzymatic nucleic acid useful for
enzymatic nucleic acid; H-Ras; N-Ras; HIV; cytostatic; anti-HIV;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 58; Page 129; 185pp; English.
                        anti-rheumatic; cancer; AIDS; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1627 GGCCCCAGCCAGCCAGCG 1643
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       BP
                                                                                                                                                                                                                                   29-MAY-2001; 2001US-0294140P.
06-JUN-2001; 2001US-0296249P.
10-SEP-2001; 2001US-0318471P.
                                                                                                                                                                                             29-MAY-2002; 2002WO-US016840.
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                                                                                                                                                                                                                                                                                                                           (RIBO-) RIBOZYME PHARM INC.
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Matches 15; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Hepatitis C virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WO200281494-A1.
                                                                                                          WO200297114-A2
                                                                    Homo sapiens.
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                                                                                                                                                                                                                                                                                                                                                                     Mcswiggen J;
                                                                                                                                                        05-DEC-2002.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The invention relates to a novel short interfering RNA (siRNA) nucleic acid molecule, that modulates expression of a nucleic acid molecule encoding HER2, K-Ras, H-Ras, N-Ras, human immunodeficiency virus (HIV) or a component of HIV. The nucleic acid molecule in anti-HIV, and anti-HIV, and molicic activity. The nucleic acid molecules are useful for reducing HER2, K-Ras, H-Ras, and HIV activity in a cell. The nucleic acid anservatic cartivity or activity in a cell. The nucleic acid anservatic activity in a molecule are also useful for treating breast, ovarian, colorectal, lung, prostate, also useful for treating breast, ovarian, colorectal, lung, prostate, shown in ABZ59889 - ABZ62216, ABZ64544 - ABZ65531, ABZ66520 - ABZ65244, ABZ65530 - ABZ65531, ABZ652310, ABZ
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Novel short interfering RNA and enzymatic nucleic acid useful for treating cancer, modulates the expression of a nucleic acid encoding HBR2, K-Ras, H-Ras, N-Ras, and human deficiency virus sequences.
                                                                                                                                                                                                               Human; ribozyme; short interfering RNA; siRNA; HER2; K-Ras;
enzymatic nucleic acid; H-Ras; N-Ras; HIV; cytostatic; anti-HIV;
anti-rheumatic; cancer; AIDS; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human; ribozyme; short interfering RNA; siRNA; HER2; K-Ras;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 17 BP; 6 A; 5 C; 4 G; 0 T; 2 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 4; Page 143; 185pp; English
                                                                                                                                                                                  Human HER2 DNAzyme substrate #557
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human H-Ras DNAzyme target #850.
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                                                     ABZ65100 standard; RNA; 17 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             29-MAY-2001; 2001US-0294140P.
06-JUN-2001; 2001US-0296249P.
10-SEP-2001; 2001US-0318471P.
                                                                                                                                                                                                                                                                                                                                                                                                                                        29-MAY-2002; 2002WQ-US016840
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ABZ62059 standard; RNA; 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (RIBO-) RIBOZYME PHARM INC.
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                                                                                                                                        (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2003-140484/13.
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                        WO200297114-A2
                                                                                                                                                                                                                                                                                                                Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mcswiggen J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   21-MAR-2003
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                                                                                                                                        21-MAR-2003
                                                                                                                                                                                                                                                                                                                                                                                                05-DEC-2002
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                                                                                              ABZ65100;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          RESULT 1349
        RESULT 1348
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ABZ62059/
                                 ABZ65100
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               amberzyme, G-cleaver ribozyme, decoy molecule, aptamer;
HBV reverse transcriptase; Enhancer I region; viral replication;
degemerative, disease state, HBV infection; HCV infection; cirrhosis;
liver failure; hepatocellular carcinoma; hepatotropic; cytostatic;
virucide; antiinflammatory; substrate; ss.
                                                                                                                                                         Gaps
                                                                                                                                                         0
                                                                    0.8%; Score 13.8; DB 1; Length 17;
88.2%; Pred. No. 9e+02;
tive 0; Mismatches 2; Indels
Sequence 17 BP; 1 A; 6 C; 7 G; 0 T; 3 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              HCV DNAzyme substrate sequence #1574.
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RNA stability; RNA expression; RNA synthesis; antisense; enzymatic nucleic acid; hammerhead riboxyme; DNAzyme; inozyme; denzymatic nucleic acid; hammerhead riboxyme; drowsyme; decoy molecule; aptamer; HBV reverse transcriptase; Enhancer I region; viral replication; degenerative; disease state; HBV infection; HCV infection; cirrhosis; liver failure; hepatocellular carcinga; hepatotropic; cytostatic; virucide; antiinflammatory; substrate; ss.

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Nucleic acid molecule; Hepatitis C virus; HCV; Hepatitis B virus; HBV;
                                                                                                                                                                                                                                                                                                                                                                                                    0.8%; Score 13.8; DB 1; Length 17; 76.5%; Pred. No. 9e+02; Attive 2; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   HCV DNAzyme substrate sequence #652.
                                                                                                                                                                                                                          Claim 1; Page 262; 387pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                  351 GGGGTCTGATGGGGAGA 367
                                                                                                                                                                                                                                                                                                                                                                                                                                          1 GGGUCUGGCGGGGAGA 17
                                        08-JUN-2001; 2001US-00877478
08-JUN-2001; 2001US-0296876P
24-OCT-2001; 2001US-0335059P
05-DEC-2001; 2001US-0337055P
                    26-MAR-2002; 2002WO-US009187.
                                                                                RIBOZYME PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ACD58066 standard; RNA; 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                        Macejak D,
Roberts E;
                                                                                                     MCSWIGGEN J. MORRISSEY D.
                                                                                                                                                                              WPI; 2003-229207/22.
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Best Local Similarity
                                                                                              MACEJAK D.
                                                                                                                           LEE P.
DRAPER K.
                                                                                                                                          ROBERTS E.
                                                                                                                   PAVCO P.
                                                                                       BLATT
                                  26-MAR-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    23-SEP-2003
      17-0CT-2002
                                                                                                                                                        Blatt L, N
Draper K,
                                                                                                                                                                                                                                                                                                                                                                                                                    13;
                                                                                                                                                                                                            infection.
                                                                                                                                                                                                                                                                                                                                                                         invention
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                                                                                                                   (PAVC/)
(LEEP/)
(DRAP/)
                                                                                                                                          (ROBE/)
                                                                                (RIBO-)
                                                                                      BLAT/)
                                                                                               MACE/)
                                                                                                     MCSW/)
                                                                                                            (MORR/)
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Matches
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The present invention relates to nucleic acid molecules which modulate the synthesis, expression and/or stability of Hepatitis C virus (HCV) or hepatitis B virus (HBV) RNA. The nucleic acid molecules include antisense and enzymatic nucleic acids such as hammerhead ribozymes. DNAzymes, inozymes, zinzymes, amberzymes, and G-cleaver ribozymes. Also disclosed are nucleic acid decoy molecules and aptamers that bind to HBV reverse transcriptuse and/or HBV reverse transcriptuse and/or HBV reverse transcriptuse primer sequences, as well as oligonuclectides that specifically bind the Enhancer I region of HBV DNA. The nucleic acids may be used to modulate the expression of HBV compounds and/or potential therapies disclosed is a method for screening compounds and/or potential therapies disclosed is a method for screening compounds that modulate the expression and/or replication of HCV. The compounds that modulate the expression and/or replication of HCV. The compounds methods of the invention are useful for the treatment of degenerative and disease states related to HBV and HCV infection, replication and gene expression such as cirrhosis, liver failure, and hepatocellular carcinoma. The present sequence represents a substrate for one of the HCV invention. Novel compound useful for treating cirrhosis, liver failure, hepatocellular carcinoma, or condition associated with hepatitis C virus Lee P; Mcswiggen J, Morrissey D, Pavco P, Sequence 17 BP; 2 A; 2 C; 11 G; 0 T; 2 U; 0 Other;

08-JUN-2001; 2001US-00877478. 08-JUN-2001; 2001US-0296876P. 24-OCTJ-2001; 2001US-0335059P. 05-DEC-2001; 2001US-0337055P.

RIBOZYME PHARM INC

MACEJAK D. MCSWIGGEN J. MORRISSEY D.

(BLAT/) (MACE/) (MCSW/) (RIBO-)

LEE P. DRAPER K. ROBERTS E.

(MORR/) (PAVC/) (LEEP/) (DRAP/) (ROBE/)

PAVCO P.

26-MAR-2002; 2002WO-US009187

Hepatitis C virus

WO200281494-A1.

17-0CT-2002

2001US-00817879

26-MAR-2001;

ь; Lee

Pavco P,

Mcswiggen J, Morrissey D,

Macejak D, Roberts E;

Blatt L, I Draper K,

WPI; 2003-229207/22.

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The present invention relates to nucleic acid molecules which modulate the synthesis, expression and/or stability of Hepatitis C virus (HCV) or Hepatitis B virus (HBV) RNA. The nucleic acid molecules include antisense and enzymatic nucleic acids such as hammerhead ribozymes, DNAzymes, include antisense includes antisenses and enzymatic nucleic acids such as hammerhead ribozymes. DNAzymes, includes and aptamers that bind to HBV reverse transcriptase primer sequences, as well transcriptase and/or HBV reverse transcriptase primer sequences, as well as oligonucleotides that specifically bind the Enhancer I region of HBV DNA. The nucleic acids may be used to modulate the expression of HBV DNA. The nucleic acids may be used to modulate the expression of HBV compounds and/or potential therapies directed against HBV, and compounds that modulate the expression and/or replication of HCV. The compounds and methods of the invention are useful for the treatment of degenerative and disease states related to HBV and HCV infection, replication and gene expression such as cirrhosis, liver failure, and hepatocellular carcinoma. The present sequence represents a substrate for one of the HCV infection and present
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Novel compound useful for treating cirrhosis, liver failure, hepatocellular carcinoma, or condition associated with hepatitis C virus
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Pred. No. 9e+02;
0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 17 BP; 2 A; 7 C; 3 G; 0 T; 5 U; 0 Other;
                                                                                                                                                                      Claim 1; Page 245; 387pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1432 GCAGAGGATGCCATGAA 1448
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nes 15; Conservative
                                                                                       infection.
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Mus musculus.
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04-DEC-2003
                                                                                                       27-MAR-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ACC68806), which are associated with tumour suppression, tumour reversion, apoptosis and virus resistance. The oligomuclectides are useful as (1) as probes and primers for detecting, identifying quantifying and/or amplifying nucleic acid, e.g. as one component of gene chip, in vitro as (anti) sense reagents; and (2) for production of recombinant polypeptides. The oligomuclectides are useful for preparation of pharmaceuticals for prevention and/or treatment of viral diseases that are characterised by development of tumours or cell degeneration, specifically cancer but also Alzheimer's disease and schizophrenia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New isolated nucleic acid, useful for treating viral diseases associated with tumors and cell degeneration, also related polypeptides, antibodies and transfected cells.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Cytostatic; virucide; neuroprotective; nootropic; neuroleptic; murine; tumour suppression; tumour reversion; apoptosis; virus resistance; viral disease; tumour; cell degeneration; cancer; Alzheimer's disease;
                                                                                                                                                                                                                Cytostatic, virucide, neuroprotective, nootropic, neuroleptic, murine, tumour suppression, tumour reversion, apoptosis, virus resistance, viral disease, tumour, cell degeneration, cancer, Alzheimer's disease,
                                                                                                                                                                         Murine oligonucleotide associated with tumour supression, SEQ ID 5972.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Murine oligonucleotide associated with tumour supression, SEQ ID 5678
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The present invention relates to murine oligonucleotides (ACC62754-
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 17 BP; 2 A; 11 C; 2 G; 2 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Tuijnder M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1482
                                         ACC68725 standard; DNA; 17 BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       17-SEP-2001; 2001FR-00011979.
                                                                                                                                                                                                                                                                                                                                                                                                                                                          17-SEP-2002; 2002WO-IB004210.
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                                                                                                                             (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2003-333167/31.
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                                                                                                                                                                                                                                                                                     schizophrenia; ss.
                                                                                                                                                                                                                                                                                                                                                                     WO2003025176-A2.
                                                                                                                                                                                                                                                                                                                            Mus musculus
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       01-JUL-2003
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                  ACC68725,
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                                                           MAKAXEX DX SX DX S
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New isolated nucleic acid, useful for treating viral diseases associated with tumors and cell degeneration, also related polypeptides, antibodies and transfected cells.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            present invention relates to murine oligonucleotides (ACC62754-
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38.2%; Pred. No. 9e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 17 BP; 5 A; 7 C; 2 G; 3 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Disclosure; Page 694; 738pp; French
                                                                                                                                                                                                                                                                                                                                                            Tuijnder M;
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                                                                                                                                                                                                                    17-SEP-2001; 2001FR-00011979.
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                                                                                                                                            17-SEP-2002; 2002WO-IB004210
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                                                                                                                                                                                                                                                                                                                                                            Telerman A, Amson R,
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WO2003025176-A2.
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Tuijnder M;

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New nucleic acid encoding human prostate membrane-specific antigen, useful e.g. for treatment of tumors and viral infection, also related polypeptide and antibodies.
                                                   Disclosure; Page 366; 771pp; French.
Amson R,
            WPI; 2003-441574/41.
 Telerman A,
                                                                                                                                                                                  potentially
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The invention relates to the isolation of 6327 nucleotide sequences, fragments of at least 15 consecutive nucleotides of these nucleotides, a sequence having at least 80% identity, after optimal alignment, with the nucleotides, or the complement, or corresponding RNA, of the nucleotides. The nucleotides are used as probes or primers for detecting, identifying quantifying and/or amplifying nucleic acids, as in vitro sense and antisense sequences, of incleotides involved in tumour suppression or reversion, apoptosis and or viral resistance, to produce recombinant polypeptides, and to prepare transgenic animals, as recombinant models. The nucleotides (also vectors containing them and cells containing the vectors), the encoded polypeptides and antibodies (Ab) against the polypeptide are useful for prevention and/or treatment of tumours or diseases characterized by development of tumours or cell degeneration (e.g. Alzheimer's disease or schizophrenia). Analysis of the expression of the nucleotides can be used for diagnosis and/or prognosis of these diseases. The nucleotides and polypeptides can also be used to screen for their specific interactive molecules, useful for treating diseases associated with abnormal expression of the nucleotides.

. 0 0.8%; Score 13.8; DB 1; Length 17; 88.2%; Pred. No. 9e+02; 2; Indels Sequence 17 BP; 7 A; 2 C; 6 G; 2 T; 0 U; 0 Other; 0; Mismatches 127 GATCGGATGAAGAT 143 17 GATCGGAAGCAGAAGAT 15; Conservative Best Local Similarity Query Match Matches g à

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Gaps

ADC03574 standard; DNA; 17 BP. 18-DEC-2003 (first entry) Homo sapiens ADC03574; RESULT 1355

Human Na/H exchanger-like protein 1 gene oligonucleotide #21

sodium/hydrogen exchanger like protein; ss; gene therapy; vaccine; sodium/hydrogen exchanger lik NHELP1; passive replacement therapy; vaccine; diagnosis.

EP1273660-A2

08-JAN-2003

25-JAN-2002; 2002EP-00001160. 30-JAN-2001; 2001WO-US000666. 23-MAY-2001; 2001US-00864761. 21-DEC-2001; 2001US-0343331P.

(AEOM-) AEOMICA INC

Gu Y;

WPI; 2003-302724/30

. 0 The invention relates to a nucleic acid molecule which encodes a Na+/H+ exchanger like protein (NHELP1). The NHELP1 nucleic acid molecule, NHELP1 polypeptide, an antibody against the protein or its antigen-binding fragment is useful in therapy. The NHELP1 nucleic acid molecule, NHELP1 polypeptide and an agonist are particularly useful for manufacturing a medicament for treating or preventing a disorder associated with decreased expression or activity of human NHELP1. The antibody or its antigen-binding fragment, and an antagonist, are useful for manufacturing a medicament for treating or preventing a disorder associated with increased expression or activity of human NHELP1. The NHELP1 uncleic acid or protein is useful as passive replacement therapy, as a vaccine, or in diagnostic methods. This sequence corresponds to a 17-mer oligonucleotide spanning the sequence of the human NHELP1 gene (ADC03514). we human sodium-hydrogen exchanger like protein 1 (NHELP1), useful as assive replacement therapy or as a vaccine for treating or preventing isorders associated with aberrant expression or activity of human Gaps . 0 0.8%; Score 13.8; DB 1; Length 17; 38.2%; Pred. No. 9e+02; ive 0; Mismatches 2; Indels Sequence 17 BP; 7 A; 4 C; 2 G; 4 T; 0 U; 0 Other; Example 2; SEQ ID NO 61; 468pp; English 1251 TATCTTAGGAACCCCAA 1267 1 TATCTAAGGAATCCCAA 17 88.2%; Conservative Local Similarity 15; Query Match NHELP1 Matches

Human tumour suppression/reversion-related DNA sequence SeqID1138. ADI48635 standard; DNA; 17 BP. (first entry) 15-APR-2004 ADI48635; RESULT 1356 ADI48635/c

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cytostatic; virucide; neuroprotective; nootropic; neuroleptic; probe; primer; PCR; gene chip; antisense; viral disease; tumour; cell degeneration; cancer; Alzheimer's disease; schizophrenia; ds; human. tumour suppression; tumour reversion; apoptosis; virus resistance;

sapiens Homo

WO2003025177-A2.

27-MAR-2003

17-SEP-2001; 2001FR-00011980. 17-SEP-2002; 2002WO-IB004523.

(MOLE-) MOLECULAR ENGINES LAB

Tuijnder M;

Amson R,

Telerman A,

WPI; 2003-313354/30.

New isolated nucleic acid, useful for treating viral diseases associated with tumors and cell degeneration, also related polypeptides, antibodies and transfected cells.

Disclosure; SEQ ID NO 1138; 30pp; French.

This invention relates to novel isolated nucleic acid sequences involved in the phenomena of tumour suppression, tumour reversion, apoptosis and/or resistance to viruses. The invention may be useful for the development of compounds with a cytostatic, virucide, neuroprotective,

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probes and primers for detecting, indentifying, quantifying and/or amplifying nucleic acid, for example as one component of a gene chip, in vitro as antisense reagents and for production of recombinant polypeptides. The invention may therefore be useful for preparation of preparated for prevention and/or treatment of viral diseases that are characterised by development of tumours or cell degeneration, specifically cancer but also Alzheimer's disease and schizophrenia. The present sequence is that of a nucleic acid sequence of the invention. Note: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO
                                                                                                                                                                                                                                                                                                                      ftp.wipo.int/pub/publishedpct_sequences
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Score 13.8; DB 1; Length 17;
Pred. No. 9e+02;
0; Mismatches 2; Indels
Sequence 17 BP; 4 A; 3 C; 6 G; 4 T; 0 U; 0 Other;
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                                                                         195 CAATGGTGCCCCTGAGC
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Best Local Similarity 88.2.
Local Si Conservative
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Gaps

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Human tumour suppression/reversion-related DNA sequence SeqID2459. ADI49956 standard; DNA; 17 (first entry) 15-APR-2004 ADI49956; RESULT 1357 ADI49956,

tumour suppression; tumour reversion; apoptosis; virus resistance; cytostatic; virucide; neuroprotective; noodropic; neuroleptic; probe; primer; PCR; gene chip; antisense; viral disease; tumour; cell degeneration; cancer; Alzheimer's disease; schizophrenia; ds; human.

Homo sapiens

WO2003025177-A2.

27-MAR-2003

17-SEP-2002; 2002WO-IB004523.

17-SEP-2001; 2001FR-00011980.

(MOLE-) MOLECULAR ENGINES LAB

Telerman A, Amson R,

Tuijnder M;

WPI; 2003-313354/30.

New isolated nucleic acid, useful for treating viral diseases associated with tumors and cell degeneration, also related polypeptides, antibodies transfected cells. and

Disclosure; SEQ ID NO 2459; 30pp; French.

in the phenomena of tumour suppression, tumour reversion, aguationed intolved in the phenomena of tumour suppression, tumour reversion, apptosis and/or resistance to viruses. The invention may be useful for the development of compounds with a cytostatic, virucide, neuroprotective, nootropic or neuroleptic activity. The DNA sequences may be useful as probes and primers for detecting, indentifying, quantifying and/or amplifying nucleic acid, for example as one component of a gene chip, in vitro as antisense reagents and for production of recombinant polypeptides. The invention may therefore be useful for preparation of phramaceuticals for prevention and/or treatment of viral diseases that are characterised by development of tumours or cell degeneration, specifically cancer but also Alzheimer's disease and schizophrenia. The This invention relates to novel isolated nucleic acid sequences involved

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      present sequence is that of a nucleic acid sequence of the invention.

Note: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      antisense oligonucleotide, neurite growth inhibitor; NOGO; prostaglandin D2 receptor; PTGDR; IkappaB kinase; IKK; protein kinase PKR; cerebrovascular accident; central nervous system injury; CNS injury; spinal cord injury; cancer; melanoma; lymphoma; glioma; inflammatory disease; rheumatoid arthritis; restenosis; asthma; Crohn's disease; diabetes; obesity; autoimmune disease; lupus; multiple sclerosis; transplant rejection; graft rejection; ischaemia; reperfusion; glomerulonephritis; sepsis; allergy; asthma; allergic rhinitis; atopic dermatitis; human PTGDR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The invention comprises nucleic acids (e.g. antisense oligonucleotides) that down regulate the expression or inhibit the function of a receptor for a neurite growth inhibitor, NOGO, prostaglandin Dz receptor (PTGDR), IkappaB kinase (IKK), or protein kinase PKR. The nucleic acids of the invention are useful for treating: cerebrovascular accident, central nervous system (CNS) injury, spinal cord injury, cancer (e.g. melanoma, lymphoma or glioma), inflammatory disease (e.g. rheumatoid arthritis, restenosis or asthma), croin's disease, diabetes, obesity, autoimmune disease, lupus, multiple sclerosis, transplant/graft rejection, ischaemia/repetusion injury, glomerulonephritis, sepsis, and allergic conditions (e.g. asthma, allergic rhinitis or atopic dermatitis). The nucleic acids of the invention are also useful for down-regulating the
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sequence of the invention.
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                                                                                                                                    Length 17;
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                                                                                                                                                                           2; Indels
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                                                                                              Sequence 17 BP; 3 A; 5 C; 2 G; 7 T; 0 U; 0 Other;
                                                                                                                                      8; DB 1;
9e+02;
                                                                                                                                      Score 13.8; DB
Pred. No. 9e+02
0; Mismatches
  sequence is that of a nucleic acid
                                                            at ftp.wipo.int/pub/publishedpct_sequences
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human PTGDR substrate sequence #1013.
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29-MAY-2001; 2001US-0294412P.
28-AUG-2001; 2001US-0315315P.
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                                                                                                                                        0.8%;
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                                                                                                                                                                                                                                                           AGGGTAAAATCGGATC
                                                                                                                                                                                                                                                                                                                                                      standard; RNA; 17
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                                                                                                                                                                               15; Conservative
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                                                                                                                                                          Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     substrate; ds
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Best Local
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expression of a target gene and as a diagnostic tool to examine genetic drifts and mutations within diseased cells or to detect the presence of target RNA in a cell. The present RNA sequence represents a human PKR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     antisense oligonucleotide; neurite growth inhibitor; NOGO; prostaglandin D2 receptor; PTGDR; IkappaB kinase; IKK; protein kinase PKR; cerebrovascular accident; central nervous system injury; CNS injury; spinal cord injury; cancer; melanoma; lymphoma; glioma; inflammatory disease; rheumatoid arthritis; restenosis; asthma; Crohn's disease; diabetes; obesity; autoimmune disease; lupus; multiple sclerosis; transplant rejection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Novel enzymatic nucleic acid that down-regulates expression of neurite growth inhibitor receptor, prostaglandin D2 receptor, IkappaB kinase or protein kinase PKR genes, for treating cancer and inflammatory disease.
                                                                                                                                                                                   Gaps
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                                                                                                                                       Score 13.8; DB 1; Length 17;
Pred. No. 9e+02;
3; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human NOGO receptor zinzyme substrate sequence #86.
                                                                                                   Sequence 17 BP; 3 A; 5 C; 6 G; 0 T; 3 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NOGO receptor zinzyme; substrate; ds.
                                                                                                                                                                              3;
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                                                                                                                                           0.8%;
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29-MAY-2001; 2001US-0294412P.
28-AUG-2001; 2001US-0315315P.
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                                                                                                                                                                                                                                                                                                                                                             ADL47099 standard; RNA; 17
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                                                                                                                                                                                 Conservative
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                                                                                                                                         Query Match
Best Local Similarity
Matches 12; Conserv
                                                            substrate sequence
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expression of a target gene and as a diagnostic tool to examine genetic drifts and mutations within diseased cells or to detect the presence of target RNA in a cell. The present RNA sequence represents a human NOGO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              antisense oligonucleotide; neurite growth inhibitor; NOGO; prostaglandin D2 receptor; PTGDR; IkappaB kinase; IKK; protein kinase PKR; cerebrovascular accident; central nervous system injury; CNS injury; spinal cord injury; cancer; melanoma; lymphoma; glioma; inflammatory disease; rheumatoid arthritis; restenosis; asthma; Crohn's disease; diabetes; obeaity; autoimmune disease; lupus; multiple sclerosis; transplant rejection; graft rejection; ischaemia; reperfusion; glomerulonephritis; sepsis; allergy; asthma; allergic rhinitis; atopic dermatitis; Human IKK-gamma;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Novel enzymatic nucleic acid that down-regulates expression of neurite growth inhibitor receptor, prostaglandin D2 receptor, IkappaB kinase or protein kinase PKR genes, for treating cancer and inflammatory disease.
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                                                                                                                                       Score 13.8; DB 1; Length 17; Pred. No. 9e+02; 2; Indels 3; Mismatches 2; Indels
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                                                                                                  Sequence 17 BP; 0 A; 6 C; 7 G; 0 T; 4 U; 0 Other
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human IKK-gamma substrate sequence #484.
                                                           receptor zinzyme substrate sequence.
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29-MAY-2001; 2001US-0294412P.
28-AUG-2001; 2001US-0315315P.
                                                                                                                                       0.8%;
                                                                                                                                                                                                                                                                                                                                                          ADL47974 standard; RNA; 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (RIBO-) RIBOZYME PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                      (first entry)
                                                                                                                                                         12; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Blatt L, Chowrira B,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2003-058513/05.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  substrate; ds
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Unidentified,
                                                                                                                                                                                                                                                                                                                                                                                                                                      20-MAY-2004
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                                                                                                                                                                                                                                                                                                                                                                                               ADL47974;
                                                                                                                                       Query Match
                                                                                                                                                           Local
                                                                                                                                                                                                                                                                                                                     RESULT 1360
                                                                                                                                                           Best Loca
Matches
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Gaps

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Enzymatic nucleic acid molecules which specifically cleave RNA derived from hepatitis C virus (HCV), useful for the treatment of HCV infections,
expression of a target gene and as a diagnostic tool to examine genetic drifts and mutations within diseased cells or to detect the presence of target RNA in a cell. The present RNA sequence represents a human PKR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The invention relates to an enzymatic nucleic acid molecule which specifically cleaves RNA derived from hepatitis C virus (HCV), in which the binding arms of the enzymatic nucleic acid molecule comprises sequences complementary to any of the defined substrate sequences given in the specification. The nucleic acid molecule may be administered for the treatment of HCV infections, especially in combination with type I interferons. The present sequence represents a HCV DNAzyme substrate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ss; enzymatic nucleic acid; RNA cleavage; hepatitis C virus; HCV; HCV infection; type I interferon; DNAzyme.
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Pred. No. 9e+02;
2; Mismatches 2; Indels
                                                                                                                                                       Length 17;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               from hepatitis C virus (HCV), useful for the treatment of especially in combination with type I interferon therapy.
                                                                                                                                                                                                   2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Macejack
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 17 BP; 2 A; 2 C; 11 G; 0 T; 2 U; 0 Other;
                                                                                                                Seguence 17 BP; 3 A; 5 C; 6 G; 0 T; 3 U; 0 Other;
                                                                                                                                                         8; DB 1;
9e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Pavco PA,
                                                                                                                                                                                                       Mismatches
                                                                                                                                                            Score 13.8;
Pred. No. 9e
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 1; SEQ ID NO 1574; 198pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      HCV DNAzyme substrate sequence #1574.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Roberts E,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                5
                                                                                                                                                                                                     3,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            351 GGGGTCTGATGGGGAGA 367
                                                                                                                                                                                                                                                318 TGCACCAGAGATTGTGC 334
                                                                                                                                                                                                                                                                                                                                                                                                        BP.
                                                                                                                                                                                                                                                                       GGGGUCUGGCGGGGAGA 17
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                                                                                                                                                            0.8%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              18-DEC-2000; 2000US-00740332
                                                                                                                                  Query Match
Best Local Similarity /v..
Best Local Similarity /v..
                                                                                                                                                                                                                                                                                                                                                                                                        ADI84328 standard; RNA; 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mcswiggen J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       BLATT L.
MCSWIGGEN J.
ROBERTS E.
PAVCO P A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2004-031273/03.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MACEJACK D.
                                                                         substrate sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Hepatitis C virus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             US2003125270-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             03-JUN-2004
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                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADI84328;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (ROBE/)
(PAVC/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (BLAT/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (MACE/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (MCSM/)
                                                                                                                                                                                                                                                                                                                                                               RESULT 1362
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Blatt
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The invention comprises nucleic acids (e.g. antisense oligonucleotides) that down regulate the expression or inhibit the function of a receptor for a neurite growth inhibitor, NOGO, prostaglandin D2 receptor (PTGDR), IkappaB kinase (IKK), or protein kinase PKR. The nucleic acids of the invention are useful for treating: cerebrovascular accident, central
         expression of a target gene and as a diagnostic tool to examine genetic drifts and mutations within diseased cells or to detect the presence of target RNA in a cell. The present RNA sequence represents a human IKK-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Novel enzymatic nucleic acid that down-regulates expression of neurite growth inhibitor receptor, prostaglandin D2 receptor, IkappaB kinase or protein kinase PKR genes, for treating cancer and inflammatory disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         rheumatoid arthritis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   nervous system (CNS) injury, spinal cord injury, cancer (e.g. melanoma, lymphoma or glioma), inflammatory disease (e.g. rheumatoid arthritis, restenosis or asthma), Crohn's disease, diabetes, obesity, autoimmune disease, lupus, multiple sclerosis, transplant/graft rejection, ischaemia/reperfusion injury, glomerulonephritis, sepsis, and allergic conditions (e.g. asthma, allergic rhinitis or atopic dermatitis). The nucleic acids of the invention are also useful for down-regulating the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             antisense oligonucleotide; neurite growth inhibitor; NOGO; prostaglandin D2 receptor; PTGDR; IkappaB Kinase; IKK; protein kinase PKR; cerebrovascular accident; central nervous system injury; CNS injury; spinal cord injury; cancer; melanoma; lymphoma; glioma; inflammatory disease; rheumatoid arthritis.
                                                                                                                                                                                                       Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             restenosis; asthma; Crohn's disease; diabetes; obesity; autoimmune disease; Lupus; multiple sclerosis; transplant rejection; graft rejection; ischaemia; reperfusion; glomearilonephritis; sepsis; allergy; asthma; allergic rhinitis; atopic dermatitis; human PTGDR;
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                                                                                                                                                         Query Match 0.8%; Score 13.8; DB 1; Length 17; Best Local Similarity 88.2%; Pred. No. 9e+02; Matches 15; Conservative 0; Mismatches 2; Indels
                                                                                                                0 T; 4 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mcswiggen J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 161; SEQ ID NO 5428; 317pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human PTGDR substrate sequence #1014.
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                                                                                                                  Sequence 17 BP; 1 A; 11 C; 1 G;
                                                                                                                                                                                                                                                     30 GCAGAGGTAGGCAGGAG 46
                                                                                                                                                                                                                                                                                                                                                                                                          ADL51895 standard; RNA; 17 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     05-APR-2001; 2001US-00827395.
29-MAY-2001; 2001US-0294412P.
28-AUG-2001; 2001US-0315315P.
                                                                                                                                                                                                                                                                                              17 GGAGAGGTAGGCAGGG 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (RIBO-) RIBOZYME PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
                                                                           gamma substrate sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Chowrira B,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2003-058513/05.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Unidentified
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     17-0CT-2002.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         substrate;
                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADL51895;
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                                                                                                                                                                                                                                                                                                                                                                 RESULT 1361
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Gaps

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Enzymatic nucleic acid molecules which specifically cleave RNA derived from hepatitis C virus (HCV), useful for the treatment of HCV infections, especially in combination with type I interferon therapy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      breast cancer; cytostatic; gene therapy; human;
intercellular adhesion molecule; ICAM-1; human rhinovirus receptor; BB2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The invention relates to an enzymatic nucleic acid molecule which specifically cleaves RNA derived from hepatitis C virus (HCV), in which the binding arms of the enzymatic nucleic acid molecule comprises sequences complementary to any of the defined substrate sequences given in the specification. The nucleic acid molecule may be administered for the treatment of HCV infections, especially in combination with type I interferons. The present sequence represents a HCV DNAzyme substrate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
                                                                                                                                         enzymatic nucleic acid; RNA cleavage; hepatitis C virus; HCV; infection; type I interferon; DNAzyme.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Extend primer 89 used to genotype human ICAM-1/ICAM-4/ICAM-5 SNP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0.8%; Score 13.8; DB 1; Length 17; 88.2%; Pred. No. 9e+02; ative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                           Macejack D;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 17 BP; 2 A; 8 C; 2 G; 0 T; 5 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                           Roberts E, Pavco PA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 1; SEQ ID NO 652; 198pp; English.
                                                                                                               DNAzyme substrate sequence #652.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1432 GCAGAGGATGCCATGAA 1448
                              BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADP46413 standard; DNA; 17 BP.
                                                                                                                                                                                                                                                                  18-DEC-2000; 2000US-00740332.
                                                                                                                                                                                                                                                                                             18-DEC-2000; 2000US-00740332.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GGAGAGGATGCCATGGA 1
                            ADI83406 standard; RNA; 17
                                                                                  (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                          Mcswiggen J,
                                                                                                                                                                                                                                                                                                                      BLATT L.
MCSWIGGEN J.
ROBERTS E.
PAVCO P A.
                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2004-031273/03.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local Similarity
Matches 15; Conserv
                                                                                                                                                                                                                                                                                                                                                                              MACEJACK D.
                                                                                                                                                                                   Hepatitis C virus.
                                                                                                                                                                                                           US2003125270-A1.
                                                                                   03-JUN-2004
                                                                                                                                                                                                                                        03-JUL-2003
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                                                        ADI83406;
                                                                                                                                                                                                                                                                                                                                                                                                          Blatt L,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    17
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                                                                                                                                                                                                                                                                                                                                                              (PAVC/)
(MACE/)
                                                                                                                                                                                                                                                                                                                       (BLAT/)
                                                                                                                                                                                                                                                                                                                                      (MCSM/)
                                                                                                                                                                                                                                                                                                                                                   (ROBE/)
RESULT 1364
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The invention relates to a novel method for identifying a subject at risk of breast cancer comprising detecting the presence or absence of one or more polymorphic variations associated with breast cancer in a nucleic acid sample from a subject. The method of the invention has cytostatic applications and may be useful for identifying a subject at risk of breast cancer, for early diagnosis, prevention and treatment of breast cancer, possibly via gene therapy, as well as to analyse and predict a response to a breast cancer treatment and in clinical drug trials. The current sequence is that of an Extend primer (also described as probe) of the invention which was used to genotype human intercellular adhesion molecule ICAM-4/ICAM-5 gDNA. ICAM-1 (human rhinovirus receptor; BB2 (CD54;cell surface glycoprotent) P3.58) has been mapped to chromosomal position 19p13.2- (TOM-4 (Landsteiner-Wiener blood group;IW) has · 0 Identifying a subject at risk of breast cancer by detecting the presence of polymorphic variations in the ICAM, MAPKIO, KIAA0861, NUMA1 or GALE regions which are associated with breast cancer in a nucleic acid sample CD54; cell surface glycoprotein P3.58; ICAM-4; Landsteiner-Wiener blood group; ICAM-5; telencephalin; chromosome 19pl3; ss; primer; PCR; SNP; single nucleotide polymorphism; probe. G-alpha-12 inhibitor; antisense compound; cell differentiation; cancer; cell growth; metastatic growth; ss; ISIS# 20658. Gaps telencephalin) has been mapped to chromosomal position 19p13.2. .; 0 0.8%; Score 13.8; DB 1; Length 17; 88.2%; Pred. No. 9e+02; rative 0; Mismatches 2; Indels œ Reneland Seguence 17 BP; 3 A; 2 C; 9 G; 3 T; 0 U; 0 Other; Human G-alpha-12 antisense inhibitor ISIS# 20658. Kammerer SM, Example 4; Page 84; 289pp; English. Braun A, 1348 TTGAGCCACGCACCCCG 1364 BP 25-NOV-2003; 2003WO-US037948. 25-NOV-2002; 2002US-0429136P. 24-JUL-2003; 2003US-0490234P. 1670/c AAZ57670 standard; DNA; 18 17 Tráarccaccacccc (first entry) Conservative Nelson MR, (SEQU-) SEQUENOM INC. Query Match Best Local Similarity WPI; 2004-441051/41. WO2004047623-A2. from a subject. Homo sapiens 05-APR-2000 10-JUN-2004 US5998206-A. 07-DEC-1999 Roth RB, AAZ57670; RESULT 1365 AAZ57670/ g BXBXGXXXXXXXXXXX

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Gaps

Cowsert LM;

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AAT11971-84 are antisense oligonucleotides (ONS) against human cytomegalovirus (CMV) that displayed activities of at least 50 % of control (ISIS 2922 shown in AAT11961). It was found that up to 4 internal mismatches could be tolerated without loss of antiviral activity. Antisense ONS targeting CMV DNA or RNA coding for the IB1, IE2 or DNA polymerase proteins have been shown to be effective in therapy, prophylaxis and diagnosis of CMV infection. The ONS may be modified to reduce nuclease resistance and to increase their efficacy. Modifications include phosphorothioate backbones, alkyl and halogen substituted sugar field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New oligo-nucleotide inhibits cytomegalovirus replication - by binding to a portion of cytomegalovirus RNA, for the diagnosis, prophylaxis and treatment of CMV diseases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       antisense; cytomegalovirus; CMV; human; therapy; prophylaxis; diagnosis; intermediate early complex; IE1; IE2; DNA polymerase gene; ss.
                                                                           Oligomer hybridises to the transactivating protein region of the HSV genome blocking successful replication. Useful in prevention and treatment of infected cells
                                                                                                                                                                                                                                                 .
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                                                                                                                                                                                                        Length 18;
                                                                                                                                                                                                                                               Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 18 BP; 0 A; 5 C; 3 G; 10 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /note= "phosphorothioate backbone"
                                                                                                                                                                                                    Score 13.8; DB 1; Pred. No. 9.4e+02; 0; Mismatches 2;
                                                                                                                                                               Sequence 18 BP; 4 A; 3 C; 7 G; 4 T; 0 U; 0 Other
                                                                                                                                                                                                                                                 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CMV antisense oligonucleotide (ISIS 5479).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example 10; Col 17; 66pp; English.
                                        Disclosure; Fig 1; 17pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Anderson K;
                                                                                                                                                                                                                                                                                      521
                                                                                                                                                                                                                                                                                                                               17
                                                                                                                                                                                                                                                                                                                                                                                                                       .975/c
AAT11975 standard; DNA; 18 BP.
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                                                                                                                                                                                                        0.8%;
                                                                                                                                                                                                                                                                                      505 GAGGCTACCTGGAGAA
                                                                                                                                                                                                                                                                                                                                   1 GTGGGTTACCTGGAGAA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (revised)
(first entry)
                                                                                                                                                                                                                                                 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1. .18
/*tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Baker B, Draper K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 1995-292538/38.
activating protein
                                                                                                                                                                                                        Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Key
modified base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      25-JAN-1993;
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13-MAR-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      US5442049-A
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAT11975;
                                                                                                                                                                                                                                                                                                                                                                                               RESULT 1367
                                                                                                                                                                                                                                                 Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    antisense molecules. The oligonucleotides are used in modulating the function of nucleic acid molecules encoding G-alpha-12, ultimately modulating the amount of G-alpha-12 produced. The antisense compounds can be utilized for diagnostics, therapeutics, prophylaxis and as research agents and kits. They may be useful in the treatment of cancer, and
                                                                                                                                                                                                                                                                                                             a human G-alpha-12 antisense nucleotide sequence. G-alpha-12 is a
                                                                                                                                                                                                                                                                                                                               member of the G12/13 subfamily of G-proteins. The primary function of G-alpha-12 is in cell differentiation and growth. The invention relates to antisense compounds which are 8-30 noucleotides long (see AA257668-25746). The antisense molecules are targeted to the human G-alpha-12 nucleic acid molecule, and inhibit the expression of G-alpha-12. The molecules preferably have a modified internucleotide linkage, and at least one modified sugar moiety. The compounds target different regions of the human G-alpha-12 RNA. The expression of human G-alpha-12 is inhibited by contacting human cells or tissues in vitro with the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Herpes simplex virus; HSV; herpes; transactivating protein; TAP; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Inhibitor of herpes simplex virus replication - comprising oligomer complementary to initiation region of mRNA coding for HSV trans-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0.8%; Score 13.8; DB 1; Length 18; 88.2%; Pred. No. 9.4e+02; Live 0; Mismatches 2; Indels
                                                                                                                                                                                                                                Antisense inhibition of human G-alpha-12 expression.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 18 BP; 4 A; 4 C; 9 G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Herpes simplex virus replication inhibitor 294.
                                                                                                                                                                                                                                                                         Example 15; Col 38; 36pp; English
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                       99US-00256496
                                                              99US-00256496
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               555 CCTCAGCCGCCGCCTCC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAQ03964 standard; DNA; 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                                         PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (SCHE ) SCHERING CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 1990-109387/15.
                                                                                                                                                                                         WPI; 2000-095920/08.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Local Similarity
les 15; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              metastatic growth
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      30-SEP-1988;
                                                                                                         SISI (-SISI)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             26-SEP-1989;
                       23-FEB-1999;
                                                                23-FEB-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              22-AUG-1990
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18

RESULT 1366

Query Match

Matches

ð 셤 AAQ03964;

11-APR-1990 EP363059-A.

Synthetic

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                                                                                                                                                                                                                                                    /*tag= a /inote= "at least one (and preferably all) of the backbone subunits are composed of amide units, so that the oligomer consists of the nucleobases attached covalently to a polyamide backbone"
                                                                                                                                                                                                                                                                                                                                                                                                                                              New peptide nucleic acid oligomers hybridisable to cytomegalovirus or papilloma:virus - are stable anti:sense molecules with high affinity fingle stranded DNA, used for treating infections.
                    Gaps
                                                                                                                                                                                  nucleic acid; PNA; cytomegalovirus; CMV; papillomavirus;
                    0;
                                                                                                                                                                                                                                                                                                                                                                                                            Cowsert LM;
 Length 18;
                   2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                          Mirabelli CK, Ecker DJ,
                                                                                                                                                              Peptide nucleic acid targetting CMV IE2 nuc sig 2.
0.8%; Score 13.8; DB 1;
88.2%; Pred. No. 9.4e+02;
                   0; Mismatches
                                                                                                                                                                                                                                 Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 2; Page 44; 65pp; English.
                                      133 ATGAAGAAGATCAAACG 149
                                                        18 AAGAAGAAGAGCAAACG 2
                                                                                                                                                                                                                                                                                                                                                   94WO-US009039.
                                                                                                                                                                                                                                                                                                                                                                     93US-00104438,
                                                                                                      AAT01677 standard; DNA; 18
                                                                                                                                            (first entry)
                                                                                                                                                                                             antiviral; diagnostic; ss.
                   15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                          Crooke ST,
                                                                                                                                                                                                                                                                                                                                                                                        (ISIS-) ISIS PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 1995-090841/12.
          Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                          Anderson KP,
                                                                                                                                                                                                                                        misc_feature
                                                                                                                                                                                                                                                                                                                                                  09-AUG-1994;
                                                                                                                                           17-DEC-1995
                                                                                                                                                                                                                                                                                                             WO9504748-A1
                                                                                                                                                                                                              Synthetic
                                                                                                                         AAT01677;
 Query Match
                                                                                                                                                                                   peptide
                   Matches
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DNA (SEDNA) to produce antisense-type gene regulation moieties. Hence they may be used therapeutically for modulating cytomegalovirus and papillomavirus processes and also as diagnostics (e.g., as probes for specific mRNAs). PNA oligomers have high affinity for complementary single stranded DNA. They are also able to form triple helices in which a first PNA strand binds with RNA or ssDNA and a second PNA strand binds with RNA or ssDNA and a second PNA strand binds with the resulting double helix or with the first PNA strand. The PNAs possess no significant charge and are water soluble, which facilitates collular upcake. Futher, since they contain amides of non-biological amino acids, they are biostable and resistant to enzymatic degradation by proceases. The present sequence targets CNV IEZ nuclear localisation ς, New oligomers are claimed which (A) have at least one peptide nucleic acid (PNA) subunit and (B) have a sequence hybridisable to AUG region, untranslated region, intron/exon (I/E) junction or coding sequence of cytomegalovirus gene selected from DNA polymerase, IEI and IE2, or hybridisable to the E, E2, E4, E5, E6, E7, L1 or L2 reading frames of a papillomavirus. The PNAs can be used to target RNA and single stranded signal 2

Seguence 18 BP; 0 A; 5 C; 3 G; 10 T; 0 U; 0 Other;

RESULT 1370

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Nucleic acid molecule modulating VEGF receptor(s) gene expression or mRNA stability - useful for treating e.g. tumour angiogenesis, psoriasis, rheumatoid arthritis, etc., in a human patient.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The present invention describes nucleic acid molecules which modulate the synthesis, expression and/or stability of a mRNA encoding 1 or more receptors of vascular endothelial growth factor (VEGF). A patient (preferably human) having a growth factor (VEGF). A patient fms-like tyrosine kinase 1 (f1t-1), kinase insert domain containing receptor (KDR) and/or foetal liver kinase 1 (f1k-1) (e.g. tumour anglogenesis, coular diseases, postriasis and rheumatoid arthritis) can be treated by administering the nucleic acid molecule or the expression vector to the patient. AAX67275 to AAX75752 represent specific examples of nucleic acid molecules from the present invention
                                                                                                                                                                                                                                                                                                                               Vascular endothelial growth factor receptor; VBGF receptor; flt-1; flk-1; MSF; hammerhead ribozyme; hairpin ribozyme; cleavage; tumour angiogenessis; psoriasis; rheumatoid arthritis; ocular disease; fms-like tyrosine kinase 1; kinase insert domain containing receptor; foetal liver kinase 1; ss.
                                    Gaps
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 Length 18;
0.8%; Score 13.8; DB 1; Length 1
88.2%; Pred. No. 9.4e+02;
ive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                Mouse flk-1 VBGF receptor hairpin ribozyme substrate #41.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Escobedo J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 18 BP; 1 A; 6 C; 7 G; 0 T; 4 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Stinchcomb D,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 4; Page 152; 218pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1033 GACTTTGGCCTGGCCCG 1049
                                                                      133 ATGAAGAAGATCAAACG 149
                                                                                                                                                                                           BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             96US-00584040.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (RIBO-) RIBOZYME PHARM INC. (CHIR ) CHIRON CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ||||::||||:
                                                                                                      18 AAGAAGAAGAACG
                                                                                                                                                                                         AAX73494 standard; RNA; 18
                                                                                                                                                                                                                                                              (first entry)
Query Match

Best Local Similarity 88.2
Matches 15; Conservative
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Matches 12; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mcswiggen J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 1997-259017/23.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WO9715662-A2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               26-OCT-1995;
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                                                                                                                                                                                                                                                                28-JUL-1999
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                                                                                                                                                                                                                             AAX73494;
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first and second samples of components for an assay for complex formation between CP2 and a CPRE in the human IL4 promoter and causing the first sample of components to react in the assay, where the extent of complex formation between CP2 and a CPRE in the human IL4 promoter in the first assay sample is determined. A candidate affector is added to the second sample of components which is then caused to react in the assay, and the extent of complex formation between CP2 and a CPRE in the human IL4 promoter in the second assay sample is determined. The extent of complex formation between CP2 and a CPRE in the human IL4 promoter in the second assay sample is determined. The extent of complex formation between the two assay samples is compared to determine the effect of the candidate affector. The therapeutic composition comprising the affector is used for the interruption or enhancement of CP2 activity and thus regulation of Th1/Th2 cell balance, for therapeutic control of the immune response and immunological disease in a variety of conditions
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequences shown in AAV53107 to AAV53114 represent oligonucleotides homologous to the CP2 recognition element (CPRE) region and can interfere with CP27 CPRE interaction. These oligonucleotides are inhibitors of CP2 function and can be used in a therapeutic composition of the invention. A method of screening for such a CP2 function affector comprises providing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         including allergic rhinitis, allergic conjunctivitis, asthma, dermatitis, urticaria, multiple sclerosis, type I diabbetes mellitus, arthritis and parasitic infection. CP2 or dominant negative CP2 may also be useful in the management of immunodeficient disorders or malignancies by amplifying
immunological disease; allergic rhinitis, allergic conjunctivitis; CPRE; dermatitis; urticaria; multiple sclerosis; arthritis; malignancy; type I diabetes mellitus; parasitic infection; immunodeficient disorder; T helper cell response; viral antigen; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                      Use of affector(s) of CP2 function - for modulating immune responses for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                        treating e.g. allergies, auto-immune disease, infections, immunodeficiency disorders or malignancies.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 18 BP; 2 A; 6 C; 4 G; 6 T; 0 U; 0 Other;
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88.2%; Pred. No. 9.4e+02;
live 0; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                                Swendeman SL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     helper cell responses to viral antigen
                                                                                                                                                                                                                                                                                                          (JOHN-) JOHNS HOPKINS SCHOOL MEDICINE.
(SLOK ) SLOAN KETTERING INST CANCER RES.
                                                                                                                                                                                                                                                                                                                                                              Sheffery M,
                                                                                                                                                                                                                                                                                      (SCHE-) SCHEPENS EYE RES INST INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1456 TTCTTCCTCAGTCTGGG 1472
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Anti-CMV oligonucleotide #5479.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 8; Fig 4D; 58pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Trerecercagrered 17
                                                                                                                                                                                                              98WO-US003049,
                                                                                                                                                                                                                                                 97US-0037972P.
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Query Match
Best Local Similarity 88.«..,
Best Local 15, Conservative
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                                                                                                                                                                                                                                                                                                                                                              Casolaro V,
                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 1998-467194/40
                                                                                                                                                                                                              19-FEB-1998;
                                                                                                                                                                                                                                                   20-FEB-1997;
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                                                                                                   Homo sapiens
                                                                                                                                      WO9836641-A1
                                                                                                                                                                         27-AUG-1998.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAX17892;
                                                                                                                                                                                                                                                                                                                                                              Ono SJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 RESULT 1372
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAX17892/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Primers AAV47637 and AAV47638 were used to amplify the wildtype VEGF-B allele from tail DNA from F2 offspring, and can be located to exon 3 and exon 4 of the mouses VEGF-B gene. F2 mice that contain the wild-type allele were found to produce an amplified fragment of 316 bp upon PCR with these primers, however mutant alleles will not be amplified by these primers, due to most of exon 3 and all of exon 4 having been completely deleted. These mutant mice produce a non-functional protein because the deletion removes 7 out of the 8 cysteine residues, thus disrupting intramolecular disulphide bonds. The transgenic animals can be used in elucidating the effects of VEGF-B on physiological phenomena such as permeability, inflammation and/or tissue repair. (Updated on 25-MAR-2003 to correct PI field.)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            vascular
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             recognition element, IL4; promoter; asthma; therapeutic composition; function affector; Th1/Th2 cell balance regulation; immune response;
                                                                                                                                                                                           Primer, amplification, PCR, mouse, VEGF-B; allele, F2 offspring, cysteine residue, intramolecular disulphide bond, transgenic animal; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        contain cells with modified vascuuse in diagnostic and therapeutic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MHC class II Ea promoter CPRE sequence (-3 to +14 basepairs).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 13.8; DB 1; Length 18; Pred. No. 9.4e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Pekny M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Eriksson U,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 18 BP; 2 A; 5 C; 6 G; 5 T; 0 U; 0 Other;
                                                                                                                                                       Primer 1, located in exon 3 and 4 of VEGF-B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Transgenic non-human animals - which endothelial growth factor B gene for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Betsholtz C,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example 4; Page 22; 45pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                            (LUDW-) LUDWIG INST CANCER RES.
                         AAV47637 standard; DNA; 18 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                63
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAV53112 standard; DNA; 18 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1 GCCCAGCTGTGTGACTG 17
                                                                                                                                                                                                                                                                                                                                                                                  9BWO-US003212.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                       97US-0038202P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                47 GACCAGCAGTGTGACTG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
                                                                                                                      (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Aase K,
S, Li X;
                                                                                                   (revised)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 1998-457107/39
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gebre-Medhin S,
                                                                                                                                                                                                                                                                                                            WO9836052-A1
                                                                                                                                                                                                                                                                                                                                                                                  18-FEB-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                       18-FEB-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Von Euler G,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       12-NOV-1998
                                                                                                 25-MAR-2003
08-DEC-1998
                                                                                                                                                                                                                                                                                                                                             20-AUG-1998
                                                                                                                                                                                                                                                 Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAV53112;
                                                              AAV47637;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               studies
                                                                                                                                                                                                                                                                        Wus sp
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CP2
     AAV53112
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Antisense oligonucleotides (AAX17861-X17924) are targeted to a nucleic acid (AAX17925-X17948) encoding IE (immediate early) 1 or 2, or DNA polymerase of cytomegalovirus (CMV) and are able to inhibit CMV replication. Optionally the oligonucleotides include at least one 2'-(2-methoxy) sugar modification or phosphorothicate internucleotide linkages. The oligonucleotides are used to inhibit CMV infections (by in vivo or in vitro contact with cells, tissues or body fluids), especially
                                                                                                                                                                                                                                                                                                                                        New antisense oligonucleotides that target cytomegalovirus nucleic acid particularly including 2-methoxyethoxy sugar modifications, especially for treating viral retinitis, with long-lasting retention in the retina.
           Antisense; oligonucleotide; immediate early; DNA polymerase; CMV; cytomegalovirus; inhibition; replication; sugar modification; phosphorothioate; infection; retinitis; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human G-alpha-11 phosphorothioate antisense oligonucleotide #33
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Identification; genetic target; gene modulation; human; probe; antisense oligonucleotide; phosphorothicate; PCR primer; nucleotide sequence-based technology; antisense drug discovery;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0.8%; Score 13.8; DB 1; Length 18; 88.2%; Pred. No. 9.4e+02; Ative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               to treat or prevent CMV infections, particularly retinitis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 18 BP; 0 A; 5 C; 3 G; 10 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                             Chapman S;
                                                                                                                                                                                                                                                                             Anderson KP,
                                                                                                                                                                                                                                                                                                                                                                                                     Claim 7; Page 30; 99pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        133 ATGAAGAAGATCAAACG 149
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      99WO-US008268
                                                                                                                                                                                  98WO-US006895
                                                                                                                                                                                                                 97US-00838715
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAZ41129 standard; DNA; 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               26-JAN-2000 (first entry)
                                                                                                                                                                                                                                                                             Kisner DL,
                                                                                                                                                                                                                                                (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        target validation; ss
                                                                                                                                                                                                                                                                                                            WPI; 1998-568330/48.
                                                                                       Human herpesvirus 5.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity
                                                                                                                                                                                  07-APR-1998;
                                                                                                                                                                                                                 39-APR-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    13-APR-1999;
                                                                                                                      WO9845314-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WO9953101-A1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            15;
                                                                                                                                                                                                                                                                             Draper KG,
                                                                          Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Synthetic
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the expression of a target nucleic acid (tNA) sequence via binding of the compounds with the tNA sequence. The method comprises generating a library of virtual compounds in silico according to defined criteria, and evaluating in silico the binding of the virtual compounds with the tNA according to defined criteria. Also described are: (1) a method of according to defined criteria. Also described are: (1) a method of cating a set of oligonucleotides (ONS) that modulate the expression of a tNA sequence via binding of the ONS with the tNA sequence comprising generating a library of virtual compounds in silico according to defined criteria, and evaluating in silico the binding of the virtual ONS with the tNA according to defined criteria; and (2) a method of defining a set of compounds that modulate the expression of a tNA sequence via binding of the compounds with the tNA. The methods can be used for the generation and identification of synthetic compounds having defined physical, chemical or bioactive properties. Information gathered from assays of such compounds is used to identify nucleic acid sequences that are compounds is used to identify nucleic acid sequences that are compounds is used to identify nucleic acid sequences that are compounds is used to identify nucleic acid sequences that are compounds acriterials and secure and identify nucleic acid sequences that are antisense drug discovery and target validation. AA240852 to AA241220, and AA352701 to AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    .
0
                                                                                                                                                                                                                          Identifying compounds which modulate expression of nucleic acids, used to provide compounds having defined physical, chemical or bioactive properties, e.g. antisense activity.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Inhibitor-kappa B kinase-beta; IKB-beta; human; T-cell leukaemia; asthma; inflammatory response; inflammatory disease; juvenile diabetes mellitus; Graves' disease; rheumatoid arthritis; allograft rejection; diagnosis; inflammatory bowel disease; multiple sclerosis; contact dermatitis; rhintis; allergy; hyperproliferative disorder; tumour; therapy; antisense inhibitor; ss.
                                                                                                     Brooks DG;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
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                                                                                                     Sasmor HM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 18 BP; 2 A; 8 C; 3 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human IKB-Beta antisense inhibitor ISIS# 23583.
                                                                                                  Freier SM,
Vickers TA;
                                                                                                                                                                                                                                                                                                                                  Example 27; Page 108; 264pp; English.
                                                                                                  M, Baker BF, Mcneil J,
Wyatt JR, Borchers AH,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   512 ACCTGGAGAGCTGACC 528
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98US-0006763B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      88.2%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAZ31599 standard; DNA; 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Local Similarity 88.2
nes 15; Conservative
                                                  (ISIS-) ISIS PHARM INC
                                                                                                                                                                             WPI; 1999-620446/53
28-APR-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   13-JAN-2000
                                                                                                  Cowsert LM,
Obasi C, W
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              RESULT 1374
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Matches
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DP
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98US-00197008.

20-NOV-1998;

JS5977341-A

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The present sequence represents a PCR primer for Herg-3, a human erg subfamily of potassium ion channel protein. The erg genes encode potassium ion channel proteins. These proteins are implicated in the development of long Q-T syndrome, a rare, but often fatal, cardiac arrhythmia. The Herg-2 and 3 proteins can be used to identify modulators of the proteins, useful in therapeutics. The nucleic acids can be used for screening of homologues
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human G-alpha-11 phosphorothioate antisense oligonucleotide SEQ ID NO:40.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The present invention describes inhibitory antisense compounds of 8-30 nucleotides, targeted to a nucleic acid molecule encoding human G-alpha-11. AAZ19468 to AAZ19547 represent human G-alpha-11 phosphorothioate antisense oligonucleotides given in the present invention. The oligonucleotides may be useful for the treatment of diseases associated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human; G-alpha-11; antisense oligonucleotide; inhibition; expression;
                                                                                                                                                                                                                                                              Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Inhibitory antisense compounds useful for the treatment of diseases associated with G-alpha-11.
                                                                                                                                                                                                                                                              .<u>;</u>
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                                                                                                                                                                                                                    0.8%; Score 13.8; DB 1; Length 18; 88.2%; Pred. No. 9.4e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Indels
                                                                                                                                                                                                                                                              2; Indels
                                                                                                                                                                             Sequence 18 BP; 0 A; 6 C; 6 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 18 BP; 2 A; 8 C; 3 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0.8%; Score 13.8; DB 1;
88.2%; Pred. No. 9.4e+02;
iive 0; Mismatches 2;
                                                                                                                                                                                                                                                              0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 3; Col 40; 38pp; English.
                                                                                                                                                                                                                                                                                                      946
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                                                                                                                                                                                                                                                                                                                                    GCTGCTCCGTGTCCTTG 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                17 ACGTGGAGAAGGTGACC
                                                                                                                                                                                                                                                                                                      930 GCTGCTCCGTGGCCTGG
                                                                                                                                                                                                                                                                                                                                                                                                                                                AAZ19500 standard; DNA; 18
                                                                                                                                                                                                                    Query Match 0.85
Best Local Similarity 88.2
Matches 15; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                phosphorothioate; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 1999-539140/45
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Local Similarity
ses 15; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               with G-alpha-11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       04-DEC-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             04-DEC-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               15-NOV-1999
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Cowsert LM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAZ19500;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Best Loca
Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ö
                                                                                                                                                          Antisense oligonucleotides inhibiting human Inhibitor-kappa B Kinasebeta, useful for treating conditions such as inflammation, asthma, diabetes, allograft rejection, allergies, hyperproliferative disorders or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   rejection, inflammatory bowel disease, multiple sclerosis, contact demartitis, rhinitis and various allergies, or hyperproliferative disorders such as leukaemias and other tumours. (I) may also be used for detection of the above disorders
                                                                                                                                                                                                                                                                                                                                                                                                              or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Novel ion channel genes and proteins useful for identifying homologues and screening for therapeutics.
                                                                                                                                                                                                                                                                                                                       invention. (I) are 8 to 30 nucleotides in length and inhibit the expression of human Inhibitor-kappa B kinase-beta (IKB-beta). (I) inhibits the expression of human IKB-beta which plays a role in the development of T-cell leukaemia and in the activation of inflammatory responses. (I) is therefore useful for treating inflammatory diseases disorders with an inflammatory component such as asthma, juvenile diabetes mellitus, Graves' disease, rheumatoid arthritis, allograft
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human, erg subfamily; potassium ion channel protein; Herg-2; Herg-3;
cardiac arrhythmia; long Q-T syndrome; PCR primer; ss.
                                                                                                                                                                                                                                                                                                        sequence represents an antisense oligonucleotide (I) of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0.8%; Score 13.8; DB 1; Length 18; 88.2%; Pred. No. 9.4e+02; tive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 18 BP; 5 A; 5 C; 6 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human Herg-3 PCR primer SEQ ID NO:10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ..
                                                                                                                                                                                                                                                                  Claim 11; Col 40; 32pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example; Page 15; 46pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  831 CACCCTTGTCTTTGAGT 847
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    97US-00956242
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  98US-00197008
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Titus SA;
                                        (ISIS-) ISIS PHARM INC.
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                                                                                                                      WPI; 1999-619715/53.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Best Local Similarity
Matches 15; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Ganetzky BS,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WO9920760-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              21-OCT-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    22-OCT-1997;
  20-NOV-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        22-JUL-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      29-APR-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        17
                                                                                  Monia BP,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
                                                                                                                                                                                                                              tumors.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1375
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10017621-381.rng

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tumour formation; treatment; inhibit; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PCR primers AAA74956-57 were used to amplify a 316 bp fragment from exons 3 and 4 of the VEGF (vascular endothelial growth factor)-B. The primers were used to analyse VEGF-B defficient transgenic mice. VEGF-B deficient animals show heart abnormalities that appear to be caused by atrioventricular conduction defects and ischemia of the myocardium. The specification describes methods for screening a compound for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              atriorentricular conduction or ischemia modulating activity. The method comprises introducing the compound into a VEGF-B deficient non-human animal, and assaying the effect on atrioventricular conduction or ischemia. The methods are used for screening atrioventricular conduction or ischemia modulating compounds, treatment or alledviation of these conditions, diagnosis of heart disease characterized by loss of VEGF-B expression, and detecting or diagnosing VEGF-B deficiency in heart of a
                                                                                                                                VEGF-B; vascular endothelial growth factor-B; heart abnormality; ischemia; atrioventricular conduction defect; myocardium; heart disease;
                                                                                                                                                                                                                                                                                                                                                                                                 Use of vascular endothelial growth factor B deficient animals for screening atrioventricular conduction or ischemia modulating compounds, and characterization of the biological roles of the growth factor.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
                                                                                                        PCR primer used to amplify a 316 bp fragment of murine VEGF-B gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     G-alpha-i2; antisense inhibitor; infection; inflammation; prevent;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             G-alpha-i2 antisense inhibitor oligonucleotide #33 (ISIS #25844).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ..
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0.8%; Score 13.8; DB 1; Length 18;
88.2%; Pred. No. 9.4e+02;
7ative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 18 BP; 2 A; 5 C; 6 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example 4; Page 31; 58pp; English.
                                                                                                                                                                                                                                                                                                                                                  Eriksson U;
                                                                                                                                                                                                                                                                                                                       (LUDW-) LUDWIG INST CANCER RES
                         AAA74957 standard; DNA; 18 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             17
                                                                                                                                                                                                                                                                    33-MAR-2000; 2000WO-US005465.
                                                                                                                                                                                                                                                                                             99US-0160083P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GCCCAGCTGTGTGACTG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    47 GACCAGCAGTGTGACTG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAA09733 standard; DNA; 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   23-JUN-2000 (first entry)
                                                                             (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                                                                                           WPI; 2000-638114/61
                                                                                                                                                                                                                                                                                                                                                Thoren P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity
                                                                                                                                                                                                               WO200052462-A1.
                                                                                                                                                           PCR primer; ss
                                                                                                                                                                                                                                                                                           03-MAR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              test subject
                                                                            02-JAN-2001
                                                                                                                                                                                                                                         08-SEP-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           15;
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                                                   AAA74957;
                                                                                                                                                                                                                                                                                                                                                  Aase K,
                                                                                                                                                                                      wus sp
RESULT 1377
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             AAA7495
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This sequence represents an antisense oligonucleotide sequence targeted nember of the Gi subfamily of G proteins, which is involved in hormonal inhibition of adenylyl cyclase and in her regulation of plasma membrane enzymes. The expression of G-alpha-i2 has been shown to be altered in some tumnours. Mice lacking the G-alpha-i2 gene display growth retardation and develop adenocarcinoma of the colon and a form of lethal diffuse coliiis similar to ulcerative colitis in humans. The antisense molecules are useful for inhibiting the expression of G-alpha-i2 in human cells or itssues, and for treating and preventing various disorders such as
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Ribozyme; hairpin; hammerhead; gene therapy; vasotropic; restenosis; ss.
                                                                                                                                                                                                                                                                                                                                                     Novel antisense oligonucleotide containing compounds, useful for inhibiting the expression of G-alpha-i2 in human cells and tissues and treating infection, inflammation and cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    infection, inflammation and tumour formation. The antisense oligonucleotides are also useful for research and diagnostic purposes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                .;
0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Cdc 2 kinase hammerhead ribozyme recognitoin site #114.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 18 BP; 3 A; 9 C; 3 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Barber JR, Robbins JM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1636 AGGCAGCGCTGGAGGG 1652
                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 1; Col 41; 31pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match 0.8%;
Best Local Similarity 88.2%;
Matches 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 17 AGGCTGCGTCTGGAGGG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAA86683 standard; DNA; 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
                                                                                                                                                                                                                     (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                                                                              WPI; 2000-270140/23.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Tritz R, Welch PJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (IMMU-) IMMUSOL INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WO200032765-A2.
Homo sapiens.
                                                                                                                               25-JUN-1999;
                                                                                                                                                                           25-JUN-1999;
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                                         US6040179-A
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                                                                                     21-MAR-2000
                                                                                                                                                                                                                                                                    Cowsert LM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAA86683;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mammalia.
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                                                                                                                                                                                                                                                                         ņ
                                                                                                                                                                               designed to cleave RNA encoding a cyclin or cell-cycle dependent kinase other than cell-cycle dependent kinases CDK1, PCNA and Cyclin B1. Representative examples of ribozyme recognition sites are given in AAA82215 to AAA86787. The ribozyme of the invention is useful for inhibiting restenosis by introduction of the ribozyme into cells. The ribozyme is resistant to endonuclease activity and hence is efficient in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 G-alpha-12 inhibitor; antisense compound; cell differentiation; cancer; cell growth; metastatic growth; ss; ISIS# 20657.
                                             New hairpin and hammerhead ribozyme for inhibiting restenosis, cleaves RNA encoding a cyclin or cell-cycle dependent kinase other than CDKI, PCNA and Cyclin B1.
                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
                                                                                                                                                                The present invention relates to a hairpin or hammerhead ribozyme.
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                                                                                                                                                                                                                                                                                                                                                               0.8%; Score 13.8; DB 1; Length 18; 88.2%; Pred. No. 9.4e+02;
                                                                                                                                                                                                                                                                                                                                                                                                     Indels
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                                                                                                                                                                                                                                                                                                                             Sequence 18 BP; 2 A; 5 C; 5 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human G-alpha-12 antisense inhibitor ISIS# 20657
                                                                                                                                                                                                                                                                                                                                                                                                       0; Mismatches
                                                                                                                              Example 1; Page 21; 109pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Example 15; Col 38; 36pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                           1036 TTTGGCCTGGCCCGAGC 1052
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       99US-00256496
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAZ57669 standard; DNA; 18
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Best Local Similarity 88.2
Matches 15, Conservative
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                                                                                                                                                                                                                                                                                          restenosis treatment
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2000-095920/08.
                 WPI; 2000-412314/35
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SISI (-SISI)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        23-FEB-1999;
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AAZ57669/c
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part of an Escherichia coli figgellin protein except a protein expressed by E. coli H., H7, H12 or H48 type strains. The present invention also describes a method of detecting the presence of E. coli of a particular H serotype in a sample, comprising specifically hybridishing a nucleic acid, preferably at least a pair, derived from a flagellating gene, specific for a particular flagellin gene associated with the H serotype, to any coli in the sample which contain the gene, and detecting any hybridised molecules, identifying the presence of that serotype in the sample. (I) are useful for: (I) detecting the presence of E. coli of H serotype in a sample by hybridishing at least one or a pair of (I) to any E. coli in the sample and detecting the presence of E. coli by Coli in the sample and detecting the hybridised nucleic acid molecules; and (2) for detecting the presence of B. coli by the hybridishing at least one or a pair of (I) to any E. coli by hybridishing at least one or a pair of (I) to any E. coli by the sample and detecting the hybridised nucleic acid molecules; and (2) for sample and detecting the hybridised nucleic acid molecules. (I) is
                                                                                                                                                                                                                                                                                                                                                                                                    ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAZ56331 to AAZ56398 represent nucleic acid molecules (I) encoding all or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                particularly useful for detecting the combination of 0 and H antigen. Hybridised (I) when using at least one (I) is detected by southern blot analysis and, when using a pair of (I), is detected by polymerase chain reaction (PCR). AAZ56399 to AAZ56420 represent primers used in the
function of nucleic acid molecules encoding G-alpha-12, ultimately modulating the amount of G-alpha-12 produced. The antisense compounds obe utilized for diagnostics, therapeutics, prophylaxis and as research agents and kits. They may be useful in the treatment of cancer, and metastatic growth
                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Novel nucleic acid molecule useful for the detection of flagellated
bacterial strains in food, feces, etc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Escherichia coli H7 specific fliC oligonucleotide primer #1696.
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0
                                                                                                                                                                                                                                                                                                                   Length 18;
                                                                                                                                                                                                                                                                                                                                                                                                2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Flagellin, fliC, antigen, detection, PCR primer, ss.
                                                                                                                                                                                                                                          Sequence 18 BP; 2 A; 5 C; 9 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                   0.8%; Score 13.8; DB 1;
88.2%; Pred. No. 9.4e+02;
                                                                                                                                                                                                                                                                                                                                                                                                0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                             552 GCCCTCAGCCGCCGCC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             17 GACCCTCAGCCGCTGCC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
                                                                                                                                                                                                                                                                                                                   Query Match 0.83
Best Local Similarity 88.27
Matches 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SYDNEY
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Escherichia coli.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  17-MAR-2000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAZ56415;
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AAZ56415
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Sequence 18 BP; 2 A; 7 C; 7 G; 2 T; 0 U; 0 Other;

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The present invention relates to use of a DNA fragment comprising a sequence of 86 nucleotides fully defined in the specification, or its functional analogs, for regulating the expression of a gene that induces parthenocarpy in a plant, by inserting the fragment at the 5' end transcribed untranslated region of the gene. The invention is useful for transgenic plant production which do not show any malformations caused by the use of gene Deff9-iaaM in some species and cultivars, and for regulating the gene that induces parthenocarpy in a plant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Use of DNA fragment of specified length to modulate the expression of genes that induce the parthenocarpic trait in plants, by inserting the DNA fragment at the 5' end transcribed untranslated region of the gene.
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Pred. No. 9.4e+02;
0; Mismatches 2; Indels
                                                                                                             Parthenocarpy; plant; DefH9-iaaM; rolA; regulation; ss.
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                                                                                                                                                                                                                                                                                                                                                     (SPER-) IST SPERIMENTALE ORTICOLTURA. (CNDR.) CONSIGLIO NAZ DELLE RICERCHE.
                                                                                                                                                                                                                                                                                                                                                                                                             Ficcadenti N,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Disclosure, Page 11, 29pp, English.
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                                                                                                                                                                                                                                                                                                  16-JUL-1999; 99IT-RM000451.
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                                        (first entry)
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                                                                                                                                                                                                                                                                                                                                    (GINE-) GINESTRA SCARL.
                                                                                                                                                                                                                                                                                                                                                                                                             Spena A, Rotino G,
                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2001-147350/15
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Best Local Similarity
                                                                                                                                                                                    WO200105985-A1.
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                                      18-APR-2001
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                                                                                                                                                Synthetic.
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     AAF56289;
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AAF56287/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Novel antisense compounds useful for inhibiting the expression of human 3-phosphoinositide dependent protein kinase-1, useful e.g. for treating inflammation, tumors and infections.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The present sequence is one of a large number of antisense oligonucleotides which are targeted to a nucleic acid molecule encoding human 3-phosphoinositide dependent protein kinase 1 (PDK-1). The antisense compounds may be oligodeoxynucleotides or chimeric oligonucleotides, which is flanked on both sides by 2'-methoxyethyl (2'-MoDE) wings. The oligonucleotides have a phosphorothioate backbone. The antisense oligonucleotides are useful for inhibiting the expression of human PDK-1 in human cells or tissues. They are also useful for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             preventing or delaying infection, inflammation or tumours and are useful
                                      Gaps
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                                                                                                                                                                                                                                                                                                                                                   Human, PDK-1, 3-phosphoinositide dependent protein kinase-1,
antisense oligonucleotide, phosphorothioate, antiinflammatory,
cytostatic, antimicrobial, ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0.8%; Score 13.8; DB 1; Length 18; 88.2%; Pred. No. 9.4e+02; vative 0; Mismatches 2; Indels
   Length 18;
Score 13.8; DB 1; Length 1
Pred. No. 9.4e+02;
0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                   Human PDK-1 antisense oligonucleotide ISIS #29246.
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                                    0;
                                                                        1566 GCCTGACTCAGGCAGGC 1582
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AAF56289 standard; DNA; 18 BP.
                                                                                                 GCCTGACTCAGGCGGCC 18
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                                      Conservative
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Best Local Similarity
Matches 15; Conserv
 Query Match
Best Local Similarity
Matches 15; Conserv
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AAC60641/C
11D AAC606
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AC AAC606
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AAC606
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KW Human;
KW antisee
KW AC6124
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CC AC6126
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CC AC60C
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RESULT 1383

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AAF56289/c ID AAF56 XX

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is a triterpenoid responsible for paltablity to animals and resistance to pathogens and predacors. The beta-amyrin synthase encoding nucleic acid is useful for producing a transpencipality to animals and resistance to pathogens and predacors. The beta-amyrin synthase encoding nucleic acid containing it into a host cell, optionally causing or allowing recombination between the vector and the host cell genome so as to transform the host cell, and regenerating a plant from the transformed plant cell. The DNA is also useful for identifying, cloning or determining the presence of a nucleic acid in a sample and for influencing or affecting the quantity or quality of triterpenoid synthesis, preferably an oleanane type triterpene saponin synthesis, in a plant, such as altering resistance to a fungal pathogen e.g., an assonycete having a sterol-containing membrane, optionally selected from assonycete having a sterol-containing membrane, preserved from a songetted from a songette and promise transfer and promise
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                                                                                                                                                                                                                                                                                                                                                                       Gaeumannomyces graminis vars tritici and avenae, Fusarium culmorus, F. avanaeceum, Stagonospora nodorum or S. avenae, taste, palatability and/or nutritional value, of the plant, by causing or allowing expression of the DNA within the cells of the plant, following an earlier step of introducing the DNA into a cell or its ancestor. The DNA is also useful for reducing the level of triterpenoids in the plant, by causing or allowing transcription from an antisense molecule in the plant, allowing transcription from the DNA, or its part such as to reduce beta-amyrin synthase expression by co-suppression, use of a nucleic acid encoding a ribozyme specific for the DNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The invention relates to a purified polynucleotide (I) encoding a protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Novel human gene Otoferlin, underlying an autosomal recessive nonsyndromic prelingual deafness, DFNB9, and proteins encoded by the gene, implicated in deafness.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0.8%; Score 13.8; DB 1; Length 18; 88.2%; Pred. No. 9.4e+02; ive 0; Mismatches 2; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 18 BP; 4 A; 5 C; 6 G; 3 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                mouse; otoferlin; OTOF; brain;
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CENT NAT RECH SCI
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Les 15; Conservative
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(CNRS ) CNRS
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                                                                                                                                                                                                                                                                                                                                                                                                                                            The present invention relates to use of a DNA fragment comprising a sequence of 86 nucleotides fully defined in the specification, or its functional analogs, for regulating the expression of a gene that induces parthenocarpy in a plant, by inserting the fragment at the 5' end transcribed untranslated region of the gene. The invention is useful for transgenic plant production which do not show any malformations caused by the use of gene DefH9-iaam in some species and cultivars, and for regulating the gene that induces parthenocarpy in a plant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Novel beta-amyrin synthase encoding nucleic acids useful for influencing or affecting triterpene synthesis, and hence resistance to fungal pathogen, taste, palatability or nutritional value of plants.
                                                                                                                                                                                                                                                                          Use of DNA fragment of specified length to modulate the expression of genes that induce the parthenocarpic trait in plants, by inserting the DNA fragment at the 5' end transcribed untranslated region of the gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The sequence represents a PCR primer used to isolate nucleic acids
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                                                                                                                                                              Ficcadenti N, Defez R;
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                                                                       (SPER-) IST SPERIMENTALE ORTICOLTURA. (CNDR.) CONSIGLIO NAZ DELLE RICERCHE.
                                                                                                                                                                                                                                                                                                                                                                                               Disclosure; Page 11; 29pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 11; Page 63; 69pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1592 GCGTGGTGGACACCGAG 1608
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAS09667 standard; DNA; 18 BP
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16-AUG-2000; 2000GB-00020217.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Haralampidis K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GTGTGGTGGACACGGAG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                            (GINE-) GINESTRA SCARL.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2001-418055/44.
                                                                                                                                                                                                                       WPI; 2001-147350/15
                                                                                                                                                              Rotino G,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WO200146391-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Avena strigosa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          24-OCT-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         28-JUN-2001
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                                                                                                                                                              Spena A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RESULT 1385
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method for monitoring expression and/or localisation of a transgene, and production of therapeutic peptide in a mammal. The method involves quantifying or detecting the amount of marker polypeptide and/or sodium iodide symporter (NIS) linked to the product of the transgene or the transgene or the transgene or the transgene or the repeating gene released from cells into extracellular body fluid, or determining the location of labelled molecules which are transported into the cells bearing the marker peptide. The method provides convenient and effective monitoring of the level and kinetics of expression of transgenes and the tissue specific distribution of expressed transgenes in cells, tissues, animals or humans without the need for disruptive and expensive sampling methods including surgery. The transgene location can be monitored without adversely affecting the mammal or the cell. The NIS is a self protein and as such does not stimulate a host immune reaction. Furthermore, the NIS functions solely to sequester iodine into a cell, which does not adversely affect normal cellular function or overall cell
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sequence (II) encoded by a novel human gene, otoferlin (OTOF) or the long human otoferlin isoform in brain. (I) was identified as underlying an autosomal nonsyndromic prelingual deafness DFNB9, and is thus useful for detecting deafness disease in humans and for characterising the functions of proteins and genes encoding them in auditory function. AAS95622-AAS95248 represent human and mouse otoferlin coding sequences, PCR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Monitoring transgene expression and therapeutic peptide production in mammals by detecting marker polypeptides linked to transgenes or therapeutic genes released from cells into extracellular body fluid.
                                                                                                                                                                                                                                                     Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The present sequence is a self-cleaving linker. It may be used in a
                                                                                                                                                                                                                                                   0;
                                                                                                                                                                                                             Length 18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Caspase-6; protease; cleavage signal; transgene expression; transgene localisation; sodium iodide symporter; NIS; ds.
                                                                                                                                                                                                                                                 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Caspase-6 protease cleavage signal nucleotide sequence.
                                                                                                                                                                   Sequence 18 BP; 4 A; 8 C; 3 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                           Score 13.8; DB 1;
Pred. No. 9.4e+02;
                                                                                                                           primers and related sequences of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (MAYO-) MAYO FOUND MEDICAL EDUCATION & RES.
                                                                                                                                                                                                                                                   Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example 11; Page 48; 79pp; English.
                                                                                                                                                                                                                                                 0;
                                                                                                                                                                                                                                                                                       479 CACTACCAGCTGACATC 495
                                                                                                                                                                                                                                                                                                                               CACGACCAGCTGTCATC 18
                                                                                                                                                                                                                                                                                                                                                                                                                                 AAF79533 standard; DNA; 18 BP
                                                                                                                                                                                                           0.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             17-AUG-2000; 2000WO-US022566.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         99US-0149168P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          16-AUG-2000; 2000US-00639667,
16-AUG-2000; 2000US-00640198.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             29-MAY-2001 (first entry)
                                                                                                                                                                                                                                                   15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Morris J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2001-257548/26.
                                                                                                                                                                                                       Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WO200113106-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Unidentified.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     22-FEB-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAF79533;
                                                                                                                                                                                                                                                                                                                                                                                          RESULT 1387
                                                                                                                                                                                                                                                   Matches
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The present invention describes a method for treating a proliferative skin or eye disease and scarring. The method involves administering a ribozyme [1] which cleaves RNA encoding a cytckine involved in inflammation, matrix metalloproteinase (MMP), cyclin, cell-cycle dependent kinase, growth factor or a reductase, or administering a nucleic acid molecule [11] comprising a promoter operably linked to a nucleic acid molecule [11] comprising a promoter operably linked to a nucleic acid segment encoding [1]. [1] can have antipsoriatic, antisickling, charactological, cytostatic, antiseborrheic, antidiabetic, antisickling, ophthalmological, vulnerary, keratolytic and virucide activities, and cleaves RNA encoding cytokine involved in inflammation. [1] can be used in gene therapy. [1] and [11] are useful for treating proliferative skin diseases such as psoriasis, atopic dermatitis, actinic keratosis, squamous or basal cell carcinoma and viral or seborrheic wart. They can also be used for treating proliferative eye diseases such as diabetic retinopathy, vitreoretinopathy, sickle cell retinopathy, retinopathy of prematurity and retinal detachment, and for treating and preventing
                                                                                                                       ..
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                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human, ribozyme therapy, hairpin ribozyme; hammerhead ribozyme; recognition site; target; ribozyme binding site; eye disease; vulnerary; proliferative disease; skin disease; psoriasis; diabetic retinopathy; cytokine; inflammation; cell-cycle dependent kinase; cyclin; MMP; matrix metalloproteinase; growth factor; reductase; scarring; cytostatic; dermacological; antiseborrheic; antidiabetic; virucide; antisickling; ophthalmological; keratosik; keratosik; squamous cell carcinoma;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Treating proliferative skin or eye diseases and scarring, using ribozymes
                                                                                                                         Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             that cleave RNA encoding cytokines involved in inflammation, matrix metalloproteinases, growth factors and cell-cycle dependent kinases
                                                                                                                                                                                                                                                                                                                                                                                                                    Cdc 2 kinase hammerhead ribozyme recognition site SEQ ID NO:4273.
                                                                                                                       .
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  basal cell carcinoma; seborrheic wart; vitreoretinopathy; scar; sickle cell retinopathy; ss.
                                                                                  Length 18;
                                                                                                                     Indels
                                         Sequence 18 BP; 3 A; 7 C; 2 G; 6 T; 0 U; 0 Other;
                                                                              Score 13.8; DB 1;
Pred. No. 9.4e+02;
                                                                                                                     0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Disclosure; Page 385; 408pp; English.
                                                                                                                                                          1723 CATGITCACCIGCCCAC 1739
                                                                                                                                                                                                  1 CATGITCATCIGCCIAC 17
                                                                                                                                                                                                                                                                                                AAH61849 standard; DNA; 18 BP
                                                                                0.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        26-OCT-2000; 2000WO-US029500.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               99US-0161532P
                                                                                                                                                                                                                                                                                                                                                                               (first entry)
                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Tritz R;
                                                         Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (IMMU-) IMMUSOL INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2001-300427/31.
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                                                                                                                                                                                                                                                                                                                                                                               10-SEP-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               03-MAY-2001
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                                                                                                                                                                                                                                                                                                                                        AAH61849;
biology
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post-operative insertion; bone tumor; vulnerary; factor superfamily; osteopathic; gene therapy;

2000DE-01020125.

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(first entry)

BP.

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Agent for stimulating bone regrowth, useful as insert after surgery for bone cancer, comprises single sequence expressing a fusion of growth
                                                                                                                               Human cystatin C derived primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                    factor and protease inhibitor.
                                                                                                                                                       Primer; cystatin C; post-opera
transforming growth factor sup
bone regeneration; cancer; ss.
                                                                                                                                                                                                                                                                                          18-APR-2000; 2000DE-01020125.
                                                                                                                                                                                                                                                                                                                                            (UYJE ) UNIV SCHILLER JENA.
                                                                                                                                                                                                                                                                                                                                                                       Wiederanders B, Maubach G;
                                                    AAI68749 standard; DNA; 18
                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2002-018650/03.
                                                                                                                                                                                                                                       DE10020125-A1.
                                                                                                                                                                                                                                                                                                                  18-APR-2000;
                                                                                                                                                                                                               Homo sapiens
                                                                                                                                                                                                                                                                 25-OCT-2001.
                                                                                                       21-JAN-2002
                                                                              AA168749;
                           RESULT 1390
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Matches
                                       AA168749
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WAIS li. These sequences can be used in nutritional supplements, they may have cytokine, cell differentiation, cell proliferation, immunomodulatory, antiinflammatory, haematopoiesis regulating, tissue growth, chemotarctic, chaemokinetic, haemostatic, thrombolytic, tumour suppression, and tumour inhibition activities, and they may also be used in the treatment of infections, infertility, and cognitive and depressive disorders. The present sequence is a probe used to isolate the coding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     invention provides the protein and coding sequences of human
scarring such as keloid, adhesion and hypertrophic or hypertrophic burn scar. AAH57577 to AAH62099 represent sequences used in the exemplification of the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New human protein related to the ribonuclease HI large subunit, useful for treating, e.g. cancer or inflammation.
                                                                                                                                                                                                                                                                                                                             Human; clone WAIS_li; nutrition; cytckine; cell proliferation; probe;
immunomodulatory; cell differentiation; haematopoiesis; tissue growth
chemotactic; chemokinetic; thrombolytic; antiinflammatory; cancer;
cytostatic; virucide; antibacterial; fungicide; haematological;
vulnerary; contraceptive; antiinfertility; haemostatic;
tumour inhibition; ss.
                                                                                                       Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Evans C;
                                                                                                        0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length 18;
                                                                             0.8%; Score 13.8; DB 1; Length 18; 88.2%; Pred. No. 9.4e+02; tive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Lavallie ER, Collins-Racie LA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     BP; 6 A; 3 C; 8 G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 13.8; DB 1;
Pred. No. 9.4e+02;
0; Mismatches 2;
                                                     Sequence 18 BP; 2 A; 5 C; 5 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                         Human clone WA15_li coding sequence probe.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Disclosure; Page 65; 67pp; English.
                                                                                                                                  1036 TTTGGCCTGGCCCGAGC 1052
                                                                                                                                                          TTTGGCCTTGCCAGAGC 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         30-MAR-2001; 2001WO-US010246.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     31-MAR-2000; 2000US-0193769P.
                                                                                                                                                                                                                             ABA03355 standard; DNA; 18
                                                                                                                                                                                                                                                                               (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (GEMY ) GENETICS INST INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            sequence of the invention
                                                                                                         Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Treacy M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mccoy JM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity
Trahes 15; Conserve
                                                                  Query Match
Best Local Similarity
-haq 15; Conserve
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2001-639364/73.
                                                                                                                                                                                                                                                                                                                                                                                                                                                        WO200175074-A1
                                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 18
                                                                                                                                                                                                                                                                               12-FEB-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Jacobs K, N
Merberg D,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        present
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                                                                                                                                                                                                                                                      ABA03355
                                                                                                                                                                                                   RESULT 1389
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                                              This invention describes a novel agent (A) for post-operative insertion, after removal of bone tumor, comprising a nucleic acid (NA1) encoding a growth factor, especially of the transforming growth factor superfamily, linked by an oligonuclectice (NO to a sequence (NA2) encoding a protease inhibitor (Pi). The product of the invention has osteopathic and vulnerary activity and can be used for gene therapy. (A) are used to promote regeneration of bone after surgical removal of primary or metastatic bone cancers. (A) make it possible to use less extensive surgery (removal of less bone), since it reduces the risk of new metastases arising from the borders of the resected zone. It also improves growth of bone into prostheses, resulting in shorter recovery times and stronger incorporation of the prosthesis, and reduces the need for further surgery. This sequence represents a PCR primer used in the amplification of the cystatin C gene used to illustrate the method of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ..
0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0.8%; Score 13.8; DB 1; Length 18; 88.2%; Pred. No. 9.4e+02; cive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 18 BP; 1 A; 3 C; 11 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     245
Claim 8; Fig 3; 8pp; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     229 AGTGGTGGTGGTGGCGG
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Best Local S
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ID ABK1
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Conservative

2 GGACAAACTGGGGAAG 18

GGACAAACTGGGCGAGG

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Tue Nov

Page 675

probe; New nucleic acid for detecting chlorinated ethylene-decomposing bacteria used to purify soil or underground water contaminated with chlorinated Chlorinated ethylene-decomposing bacteria; 16S rRNA; 16S rDNA; ss; pPCR; primer; soil; underground water; chlorinated ethylene; KWI-De3; Chlorinated ethylene-decomposing bacteria detection DNA KWI-De3. chlorinated ethane; Dehalococcoides. Claim 1; Page 7; 22pp; English. 24-JUL-2000; 2000JP-00227580. 09-MAR-2001; 2001JP-00066001. 23-JUL-2001; 2001EP-00117844 (KURK) KURITA WATER IND LTD. WPI; 2002-173127/23. ethylene or ethane. EP1176216-A2 30-JAN-2002. Nakamura K, Synthetic.

chlorinated ethylene-decomposing bacteria (e.g. Dehalococcoides) comprising the novel nucleic acid by DNA hybridisation using the labelled probe as an indicator. The bacteria can also be detected by performing PCR using the nucleic acid as a primer and the sample nucleic acid as a template, and detecting newly synthemised DNA. A method for decomposing chlorinated ethylene or ethane comprises detecting chlorinated ethylenedecomposing bacteria using underground water or soil as a sample, and introducing the water/soil containing the bacteria, to soil or underground water contaminated by chlorinated ethylene or ethane. The methods are therefore useful for purifying soil or underground water contaminated with chlorinated ethylene or ethane. This sequence represents a nucleic acid which hybridises to nucleic acid which hybridises to nucleic acid of chlorinated ethylene-decomposing bacteria The invention relates to a nucleic acid which hybridises to the 16S ribosomal (deoxy)ribonucleic acid of chlorinated ethylene-decomposing bacteria. The nucleic acid can be used as a labelled probe for detecting

Sequence 18 BP; 5 A; 3 C; 7 G; 3 T; 0 U; 0 Other;

Gaps .. 0 0.8%; Score 13.8; DB 1; Length 18; 88.2%; Pred. No. 9.4e+02; tive 0; Mismatches 2; Indels 0, Conservative Local Similarity les 15; Conserv Query Match

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596 GCTTTGGGAAACTGGAG 612 17 GCTTCGGGAAACTGAAG

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ABS64463 standard; DNA; 18 15-NOV-2002 (first entry) ABS64463; RESULT 1392 ABS64463
IID ABS6
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DDE HUME
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XW PAUTI
XW SChi

Human TGF-beta binding PCR primer SR1 #2

Parkinson's disease; Huntington's disease; neurological disorder; schizophrenia; manic depression; mental retardation; angina pectoris; cardiovascular disease; acute heart failure; myocardial infarction; amuscular disease; muscular disease; photoreception; deafness; keratinisation disorder; catinal disease; photoreception; deafness; keratinisation disorder; cancer; ovarian cancer; melanoma; Human, NOVX; neurodegenerative disease; Alzheimer's disease; anxiety;

immune disease; diabetes; bacterial infection; fungal infection; protozoal infection; obesity; viral infection; reproductive system disorder; metabolic disturbance; anorexia; wasting disorder; chronic disease; infectious disease; dyslipidaemia, TGF-beta binding, cloning, PCR, primer, ss. disorder; inflammatory disease; WO200264791-A2 [mmunological domo sapiens

22-AUG-2002.

10-DEC-2001; 2001WO-US048369

08-DEC-2000; 2000US-0254329P. 14-DEC-2000; 2000US-025548P. 15-MAX-2001; 2001US-0291037P. 08-UJN-2001; 2001US-029773P. 08-UJN-2001; 2001US-0309288P. 29-AUG-2001; 2001US-0315639P.

01-OCT-2001; 2001US-0326393P

(CURA-) CURAGEN CORP

Alsobrook JP, Anderson DW, Burgess CE, Boldog FL, Casman SJ; Colman SD, Edinger SR, Ellerman K, Gerlach V, Gorman L, Grosse WM; Guo X, Herrmann JL, Kekuda R, Lepley DM, Li L, Macdougall JR; Millet I, Pena CEA, Peyman JA, Rastelli L, Rieger DK, Shimkets RA; Smithson G, Spytek KA, Stone DJ, Tchernev VT, Vernet CAM, Voss EZ; Zerhusen BD, Zhong H, Zhong M;

WPI; 2002-643486/69.

New NOVX polypeptides and polynucleotides useful for treating or preventing e.g. neurodegenerative diseases, neurological disorders, cardiovascular diseases and disorders, or immunological diseases.

Example 3; Page 288; 299pp; English.

medicament for treating or preventing neurodegenerative diseases (e.g. Alzheimer's disease, Parkinson's disease, or Huntington's disease),
Alzheimer's disease, Parkinson's disease, or Huntington's disease),
neurological disorders (e.g. anxiety, schizophrenia, manic depression or mental retardation), cardiovascular diseases (e.g. acute heart failure, angina pectoris or myocardial infarction), muscular diseases and diseases (including those involving photoreception, dafaness and keratinisation disorders, inflammatory and immune diseases, bacterial, fungal, protozoal and viral infections, and reproductive compounds that modulate the NOVX protein activity or expression, as well as to treat disorders characterised by insufficient or excessive production of NOVX protein forms that have decreased or aberrant activity compared to NOVX wild type protein, such as diabetes, obseity, metabolic disturbances associated with obesity, anorexia and obesity, metabolic disturbances associated with chronic diseases and various cancers, infectious diseases associated with chronic diseases and various dyslipidaemias. The nucleic acid sequences of the invention may be used in chromosome mapping, identifying an individual from minute biological samples (tissue typing), and in The present invention relates to new NOVX polypeptides. The polypeptides, polynucleotides and antibodies are useful in the manufacture of a forensic identification of a biological sample. The present nucleic acid sequence represents a cloning PCR primer that was used in the methods of the invention for amplification of the NOVX TGF-beta binding gene

Sequence 18 BP; 1 A; 10 C; 3 G; 4 T; 0 U; 0 Other;

Gaps . 0 0.8%; Score 13.8; DB 1; Length 18; 88.2%; Pred. No. 9.4e+02; Live 0; Mismatches 2; Indels ·, Local Similarity 88.2 nes 15; Conservative Query Match Best Loca Matches

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571 555 CCTCAGCCGCCGCCTCC

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                                                                                                                                                                                                                                                                                                                                                                                                                            The invention describes a new antisense compound, which is 8-30 mucleobases in length targeted to a nucleic acid molecule encoding Inhibitor-kappa B Kinase-beta that specifically hybridises with and inhibits the expression of Inhibitor-kappa B Kinase-beta. The compound is useful for preparing a composition for treating obesity, diabetes type II, inflammatory disorder or cancer e.g., leukeemia. This sequence represents an antisense oligomoulectide used to inhibit the expression of inhibitor-kappa B kinase-beta
                                                                                                                      Human; inhibitor-kappa B kinase-beta; anorectic; antidiabetic;
antiinflammatory; cytostatic; gene therapy; antisense compound; obesity;
diabetes type II; inflammatory disorder; cancer; leukaemia;
antisense oligonucleotide; ss.
                                                                                                                                                                                                                                                                                                                                                                     New antisense compound, useful for preparing a composition for treating obesity, diabetes type II, inflammatory disorder or cancer e.g.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
                                                                                                      Human Inhibitor-kappa B kinase-beta antisense oligonucleotide #12.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 18 BP; 5 A; 5 C; 6 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Beer spoilage-associated primer SEQ ID 185.
                                                                                                                                                                                                                                                                                                                                                                                                            Claim 3; Page 22; 49pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             847
                                                                                                                                                                                                                                                                                                                                 Koller
                                              ACD66643 standard; DNA; 18 BP
1 CCTCAGCGTCCGCCTCC 17
                                                                                                                                                                                                                                 24-MAY-2002; 2002US-00156610.
                                                                                                                                                                                                                                                    98US-00197008.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             831 CACCCTTGTCTTTGAGT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               caccreeccrrreagr
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DNA; 18
                                                                                     (first entry)
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Best Local Similarity 88.2
Matches 15; Conservative
                                                                                                                                                                                                                                                                                                                                Cowsert LM,
                                                                                                                                                                                                                                                                                          (MONI/) MONIA B P. (COWS/) COWSERT L M.
                                                                                                                                                                                                                                                                                                                                                  WPI; 2003-512357/48
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADE14990 standard;
                                                                                                                                                                                                                                                                                                             (KOLL/) KOLLER E.
                                                                                                                                                                                           US2003050270-A1.
                                                                                                                                                                                                                                                     20-NOV-1998;
                                                                                                                                                                                                                                                               28-JUL-1999;
                                                                                     16-SEP-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    29-JAN-2004
                                                                                                                                                                                                               13-MAR-2003
                                                                                                                                                                                                                                                                                                                                Monia BP,
                                                                                                                                                                                                                                                                                                                                                                                         Leukemia.
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                                                                  ACD66643;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RESULT 1394
                             RESULT 1393
                                      ACD66643,
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This invention describes novel oligonucleotides used in a method for detecting beer-spoilage bacteria in a sample. The bacteria detected include lattic acid bacteria of the genera Latchbacillus or Pediococcus, especially the species L. coryniformis, L. perclens, L. buchneri, L. plantarum, L. fructivorans, L. lindheri, L. casei, L. brewis or P. plantarum, or Gram-negative bacteria of the genera Pectinatus and Megasphaera, specifically P. frisingensis, P. cerevisiphilus and M. cerevisiae. The oligonucleotides of the invention provide rapid detection of spoilage bacteria (typically within 48 hours, compared with 7-12 days for conventional culture methods), can detect all relevant bacteria in parallel, can differentiate between species of the same genus, and are easy to use. ApB14806-ADB15247 represent the oligonucleotides used in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New oligonucleotides, useful for rapid detection of beer-spoilage bacteria by in situ hybridization, are specific for type, genus or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            .;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 13.8; DB 1; Length 18; Pred. No. 9.4e+02; 0; Mismatches 2; Indels
ss, primer, detection, beer-spoilage, lactic acid bacteria,
Gram-negative bacteria, spoilage bacteria.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 18 BP; 1 A; 4 C; 11 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         HLA class I allele specific primer #125.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 1; SEQ ID NO 185; 88pp; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               229 AGTGGTGGTGGTGGCGG 245
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                                                                                                                                                                                                                                                                                                                                                                              19-JUN-2001; 2001DE-01029410.
                                                                                                                                                                                                                                                                                                               19-JUN-2002; 2002WO-EP006808.
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20-DEC-2000; 2000US-00747391
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ADE13509 standard; DNA; 18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       method of the invention.
                                                                                                      Lactobacillus buchneri
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Snaidr J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2003-175243/17.
                                                                                                                                                                                                                                                                                                                                                                                                                                                 (VERM-) VERMICON AG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        US2003165884-A1.
                                                                                                                                                                            WO2002103043-A2.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Beimfohr C,
                                                                                                                                                                                                                                               27-DEC-2002
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Homo sapiens.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The present invention relates to human phenotype-associated (PA) genes (I A. ADF13307-ADF13386) which contain a Single Nucleotide Polymorphism (SNP). The SNP is given in the sequence as a variant nucleotide. Also claimed are methods for screening for agents which regulate the activity of a PA gene and reagents that modulate the activity of a PA polypeptide or a polymucleotide where the reagent is identified by the screening methods. The methods and compositions of the present invention are useful for prognosticating, diagnosing and treating cardiovascular diseases,
                                                                                                                                                 Leukocyte Antigen (HLA) genotype of a subject using hybridisation and amplification assay. The method is used for determining the HLA genotype of a subject. The present sequence represents a HLA class I allele
                                                                                                                                      The invention relates to a method of identifying a class I or II Human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             or
                                                                                                                                                                                                                                                         Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Cardiant; antiarteriosclerotic; vasotropic; cerebroprotective;
hypotensive; gene therapy; human; apolipoprotein B; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New isolated polynucleotide encoded by a phenotype-associated gene, useful for prognosticating statin therapy response, and diagnosing treating cardiovascular diseases, such as hypertension, myocardial
                                                                              Identifying class I or II Human Leukocyte Antigen genotypes using hybridization and amplification assays.
                                                                                                                                                                                                                                                                                                                                                                                                                                 Apolipoprotein E (epsilon-4 allele), BaySNP 10948, PCR primer #2.
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88.2%; Pred. No. 9.4e+02;
trive 0; Mismatches 2; Indels
                                                                                                                                                                                                           Sequence 18 BP; 3 A; 6 C; 5 G; 4 T; 0 U; 0 Other;
                                                                                                               Claim 7; SEQ ID NO 127; 66pp; English
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                                                                                                                                                                                                                                                                              503 CTGAGGGCTACCTGGAG 519
                                                                                                                                                                                                                                                                                                                                                               BP.
                                                                                                                                                                                                                                                                                                     CTGAAGCCTACCTGGAG 2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    27-FEB-2002; 2002EP-00004258
                                                                                                                                                                                                                                                                                                                                                            ADF13492 standard; DNA; 18
                                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
                                                                                                                                                                                                                                 Query Match 0.8
Best Local Similarity 88.2
Matches 15; Conservative
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            (STEM-) STEMCYTE INC
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                                                        WPI; 2003-874916/81.
                                   Tonai R;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (FARB ) BAYER AG.
                                                                                                                                                                                      specific primer.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SNP; single nucleotide polymorphism; cardiovascular associated gene; allelic variation; atherosclerosis; ischemia; reperfusion; hypertension; restenosis; arterial inflammation; myocardial infarction; stroke; primer;
such as atherosclerosis, hypertension, restenosis, arterial inflammation, myocardial infarction and stroke. The present sequence is a PCR primer, used in the examples from the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New isolated polynucleotides comprising single nucleotide polymorphisms of the cardiovascular gene, useful for assessing predisposition or susceptibility to a cardiovascular disease, e.g. atherosclerosis,
                                                                                                                                                                                                                                                                                           Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
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                                                                                                                                                                                                                0.8%; Score 13.8; DB 1; Length 18;
88.2%; Pred. No. 9.4e+02;
.ive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SNP-containing cardiovascular associated gene primer #34.
                                                                                                                                               Sequence 18 BP; 6 A; 1 C; 10 G; 1 T; 0 U; 0 Other;
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                                                                                                                                                                          Query Match
Best Local Similarity 88.2%,
ches 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                   31 CAGAGGTAGGCAGGAGG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADM92704 standard; DNA; 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match 0.8
Best Local Similarity 88.2
Matches 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Stropp U, Schwers S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         restenosis or stroke.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2003-577532/54.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (FARB ) BAYER AG.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WO2003057911-A2
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Laatikainen H;

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The invention relates to a novel method for monitoring harmful microorganisms in the paper and board industry comprising detecting the presence or absence of the harmful target microorganism by in situ hybridization using an oligonuclectide probe hybridizable with a region of a nucleic acid of the target microorganism. The method is useful for determining the need of an anti-biofilm agent in the paper or board making process, and for inhibiting the formation of agglomerates and/or biofilm, and/or removing the agglomerates and/or biofilm, and/or removing the agglomerates and/or by thermophilic microorganisms, from the surfaces of paper and board making machines. The method also enables an efficient and timesaving process for monitoring the microbiological state of a paper or board making process, and for the recognition of the microorganism before
                                                                                                                                                                                                                                                                                                                                                                       Monitoring harmful microorganisms in paper and board industry, involves detecting presence of harmful target microorganisms by in situ hybridization using oligonucleotide probe hybridizable to nucleic acid of target microorganism.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       cytomegalovirus; CMV; human, therapy; prophylaxis; diagnosis; e early complex; IE1; IE2; DNA polymerase gene; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       process failures emerge. The present sequence represents an oligonucleotide probe targeted to a region or rRNA of a target
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0.8%; Score 13.8; DB 1; Length 18; 88.2%; Pred. No. 9.4e+02;
                                                                                                                                                                                                                                                                      Rainey F, Salkinoja-Salonen M,
Vaeaetaenen P;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /note= "phosphorothioate backbone"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 18 BP; 3 A; 9 C; 3 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CMV antisense oligonucleotide (ISIS 5478).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 12; SEQ ID NO 27; 32pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              microorganism of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GCTTTGGGAAACTGGAG 612
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             N
                                                                                                                                    06-NOV-2003; 2003WO-F1000839
                                                                                                                                                                       06-NOV-2002; 2002FI-00001986
06-NOV-2002; 2002FI-00001987
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GCCTTGGGAAACTGGGG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAT11974 standard; DNA; 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1. .19
/*tag=
                                                                                                                                                                                                                                                                        Kolari M,
Vuorela P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (revised)
                                                                                                                                                                                                                                                                                                                                    WPI; 2004-419709/39.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Local Similarity
es 15; Conserv
                                                                                                                                                                                                                                   (KEMH ) KEMIRA OYJ.
                                                     WO2004042082-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Key
modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              intermediate
                Unidentified.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          25-MAR-2003
13-MAR-1996
                                                                                               21-MAY-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             antisense;
                                                                                                                                                                                                                                                                            ָט מ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         296
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
                                                                                                                                                                                                                                                                            Jurgens (Tammela
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         RESULT 1400
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Matches
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a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ð
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (HIA) genotype of a subject comprising: obtaining a sample comprising acid template nucleic acid from the subject; amplifying the template nucleic acid with HIA allele-specific forward primers and HIA allele-specific reverse primers to form amplification products; hybridising the amplification products with an HIA locus-specific capture oligonucleotide is and detecting the detectable complexes to identify the HIA genotype of the subject. The present sequence represents one of 276 HIA locus-specific capture oligonucleotides of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Identifying an HLA genotype of a subject by hybridizing the amplification products with an HLA locus-specific capture oligonucleotide and detecting the detectable complexes to identify the HLA genotype of the subject.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The invention describes a method of identifying a human leukocyte antigen
                                                                                                                                                                                                                                                                                                                    ss; primer; human leukocyte antigen; HLA; HLA genotyping; human; PCR.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
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0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ss; probe; paper; board; in situ hybridisation; anti-biofilm; agglomerate; biofilm; thermophilic microorganism.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0.8%; Score 13.8; DB 1; Length 18; 88.2%; Pred. No. 9.4e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Rubellimiccrobium rRNA oligonucleotide probe SEQ ID NO:27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 18 BP; 3 A; 6 C; 5 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                            HLA locus-specific capture oligonucleotide #125.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example 1; SEQ ID NO 127; 68pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             .
0
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                                                                                                                                                        BP.
                    47
                                                   CAGAGGGAGGAAGGAGG 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ~
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            20-DEC-2000; 2000US-00747391.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    99US-0172768P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     503 CTGAGGGCTACCTGGAG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CTGAAGCCTACCTGGAG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADO58582 standard; DNA; 18
                                                                                                                                                        ADL09359 standard; DNA; 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
                                                                                                                                                                                                                                     (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity
Thes 15; Conserva
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (STEM-) STEMCYTE INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2004-068584/07.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Tonai R;
                                                                                                                                                                                                                                                                                                                                                                                                US6670124-B1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    20-DEC-1999;
                                                                                                                                                                                                                                                                                                                                                           Homo sapiens
                                                                                                                                                                                                                                     06-MAY-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       12-AUG-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                       30-DEC-2003
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                                                                                                                                                                                                ADL09359;
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                    31
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RESULT 1399
                                                                                                                     RESULT 1398
                                                                                                                                          ADL09359
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Gaps 0

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a portion of cytomegalovirus replication - by binding to a portion of cytomegalovirus RNA, for the diagnosis, prophylaxis and treatment of CMV diseases.
                                                                                                                                                                                                                                                                                                                 AATI1971-84 are antisense oligonucleotides (ONs) against human cytomegalovirus (CMV) that displayed activities of at least 50 % of control (ISIS 2922 shown in AATI1961). It was found that up to 4 internal mismatches could be tolerated without loss of antiviral activity. Antisense ONs targeting CMV DNA or RNA coding for the IEI, IE2 or DNA polymerase proteins have been shown to be effective in therapy, polymerase proteins have been shown to be effective in therapy, prophylaxis and diagnosis of CMV infection. The ONs may be modified to reduce nuclease resistance and to increase their efficacy. Modifications include phosphorothicate backbones, alkyl and halogen-substituted sugar moieties at the 2' position. (Updated on 25-MAR-2003 to correct PF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /*tag= a //tag= a //note= "at least one (and preferably all) of the backbone
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         subunits are composed of amide units, so that the oligomer consists of the nucleobases attached covalently to a polyamide backbone"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 peptide nucleic acid, PNA; cytomegalovirus; CMV; papillomavirus;
antiviral; diagnostic; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 13.8; DB 1; Length 19;
Pred. No. 9.8e+02;
0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Peptide nucleic acid targetting CMV IE2 nuc sig 2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 19 BP; 0 A; 5 C; 4 G; 10 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Location/Qualifiers
                                                                                                                                                                                                                                                                                         Example 10; Col 17; 66pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    133 ATGAAGAAGATCAAACG 149
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAGAAGAAGAGCAAACG 2
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                                                                       93US-00009263
                                                                                                      92US-00927506
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      17-DEC-1995 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match 0.8
Best Local Similarity 88.2
Matches 15; Conservative
                                                                                                                                   (ISIS-) ISIS PHARM INC
                                                                                                                                                                 Draper K,
                                                                                                                                                                                               WPI; 1995-292538/38
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    09-AUG-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WO9504748-A1
                                                                       25-JAN-1993;
                                                                                                      19-NOV-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              nisc feature
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      16-FEB-1995
            US5442049-A
                                           15-AUG-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Synthetic
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                                                                                                                                                                   Baker B,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              field.)
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New oligomers are claimed which (A) have at least one peptide nucleic acid (PNA) subunit and (B) have a sequence hybridisable to AUG region, 5 untranslated region, intron/exon (I/E) junction or coding sequence of cytomegalovirus gene selected from DNA polymerase, IE1 and IE2, or papillomavirus. The PNAs can be used to target RNA and single stranded DNA (ssDNA) to produce antisense type gene regulation moieties. Hence they may be used therapeutically for modulating cytomegalovirus and papillomavirus processes and also as diagnostics (e.g., as probes for specific mRNAs). PNA oligomers have high affinity for complementary single stranded DNA. They are also able to form triple helices in which a first PNA strand binds with the first PNA strand binds with the first PNA strand binds with the first PNA strand. The PNAs is possess no significant charge and are water soluble, which facilitates
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0
                                                                                                                                                                                                                                                                                                                                                                                                                                                   possess no significant charge and are water soluble, which facilitates callular uptake. Futher, since they contain amides of non-biological amino acids, they are biostable and resistant to enzymatic degradation proteases. The present sequence targets CMV IE2 nuclear localisation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Salmonella typhimurium; org gene; polymerase chain reaction; PCR; primer;
                                                                                                                          papilloma:virus - are stable anti:sense molecules with high affinity for single stranded DNA, used for treating infections.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0.8%; Score 13.8; DB 1; Length 19;
88.2%; Pred. No. 9.8e+02;
.ive 0; Mismatches 2; Indels
                                                              Ecker DJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 19 BP; 0 A; 5 C; 4 G; 10 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PCR primer DP17 for org gene probe preparation.
                                                                Mirabelli CK,
                                                                                                                                                                                                 Claim 2; Page 44; 65pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          133 ATGAAGAAGATCAAACG 149
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93US-00104438
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               oxygen-regulated gene; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (GEHO ) GEN HOSPITAL CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match 0.8
Best Local Similarity 88.2
Matches 15; Conservative
                                                                Crooke ST,
                               (ISIS-) ISIS PHARM INC
                                                                                                 WPI; 1995-090841/12.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 1997-289217/26
 09-AUG-1993;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            04-AUG-1997
                                                                  Anderson KP,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 22-MAY-1997.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Miller SI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAT67044;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         signal 2
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WPI; 1998-031751/03
                                                                                               Query Match
Best Local Similarity
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Lee C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      histoplasmosis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   08-MAR-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               08-MAR-1995;
                                                                                                                                                                                                                                                                                                       01-JUN-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                              US5693501-A
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                                                                                                                                                                                                                                                                                                                                                                                                      Synthetic
                                                                                                                                                                                                                                                                             AAV01575;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     В,
                                                                                                                                                                                                                         RESULT 1404
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Jiang
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         RESULT 1405
                                                                                                                          Matches
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8 X C C C C C
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAX09121-X10268 are allele-specific oligonucleotide primers used in the isolation of various biallelic polymorphic markers found in the human genome (represented in AAX10269-X12937). These primers can be used in a method for determining polymorphic forms in an individual for use in e.g. forensics, patermity testing or for phenotypic typing for diseases such as agammaglobulinemia, adabetes inspiduas, Lesch-Nyhan syndrome, muscular dystrophy, Wiskott-Aldrich syndrome, Rabry's disease, familial hypercholesterolemia, polycystic kidney disease, hereditary spherocytosis, von Willebrand's disease, tuberous sclerosis, hereditary haemorrhagic telangiectasia, familial colonic polyposis, Ehlers-Danlos syndrome, osteogenesis imperfecta, acute intermittent porphyria, autoimmune diseases, inflammation, cancer, disease of the nervous system, infection by pathogenic microorganisms, and characteristics such as longevity, appearance (e.g. baldness, obesity), strength, speed,
New isolated Salmonella secreted proteins and related genes - used to develop products for the detection, treatment or prevention of Salmonella
                                                                               PCR primers DP15 (AAT67043) and DP17 (AAT67044) were used to amplify a 724-bp org gene probe. The probe can be used to identify the Salmonella typhimurium oxygen-regulated gene (org)
                                                                                                                                                                                                                                                                                                                                                                                                                              Polymorphism, biallelic, human; forensic; paternity testing; disease; detection; phenotypic typing; characteristic; infection; hereditary; autoimmune disease; cancer; inflammation; drug; therapy; medicament;
                                                                                                                                                                                            Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New isolated nucleic acid segments from the human genome - used for determining polymorphic forms for use in e.g. forensics, paternity testing or phenotypic typing for disease.
                                                                                                                                                                                            ö
                                                                                                                                                              0.8%; Score 13.8; DB 1; Length 19; 88.2%; Pred. No. 9.8e+02; tive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                    Human biallelic polymorphic marker downstream primer #551.
                                                                                                                                     Sequence 19 BP; 2 A; 7 C; 5 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (WHED ) WHITEHEAD INST BIOMEDICAL RES.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 16; Page 219; 310pp; English
                                                      Example 1; Page 29; 95pp; English
                                                                                                                                                                                                                      1272 GGAGACGTGGCCAGGCA 1288
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         treatment; marker; primer; ss.
                                                                                                                                                                                                                                                                                                                    AAX10245 standard; DNA; 19 BP.
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                                                                                                                                                                                                                                                GGAGAACTGGCCAGGCA
                                                                                                                                                                                                                                                                                                                                                                          (first entry)
                                                                                                                                                                 Query Match 0.8
Best Local Similarity 88.2
Matches 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Wang D,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 1998-286974/25.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WO9820165-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 05-NOV-1997;
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                                                                                                                                                                                                                                                                                                                                                                          24-MAR-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     14-MAY-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Lander ES,
                            infections
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Synthetic
                                                                                                                                                                                                                                                 18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         primers AAV01575-V01576 were used to amplify the internal transcribed spacer 1 (ITS1) sequence from the Histoplasma capsulatum large subunit ribosomal genes (AAV01567). The ITS1 sequence corresponds to the region between the 3' end of the 18S ribosomal gene and the 5' end of the 5.8S ribosomal gene. The ITS1 sequence was PCR amplified from isolated DNA from both the yeart and mycelial forms of H. capsulatum. Fragments of th sequence (e.g. AAV01568-V01574) can be used as primers and probes for H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Internal transcribed spacer, ITS; ribosomal RNA; 185; 5.8S; ss; primer; PCR; amplification; probe; hybridisation; detection; histoplasmosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Histoplasma capsulatum DNA sequences - useful as primers for diagnosing
endurance, fertility, and susceptibility or receptivity to particular drugs or therapeutic treatments. The isolated polymorphic nucleic acid segments can also be used to produce medicaments for the treatment or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
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                                                                                                                                                                                                                   Length 19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 19;
                                                                                                                                                                                                                                                                               Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  capsulatum infection (histoplasmosis) in a patient
                                                                                                                                                                                                                   Score 13.8; DB 1;
Pred. No. 9.8e+02;
0; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 19 BP; 5 A; 4 C; 5 G; 5 T; 0 U; 0 Other;
                                                                                                                                                          Seguence 19 BP; 8 A; 6 C; 4 G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 13.8; DB 1;
Pred. No. 9.8e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (INDV ) UNIV INDIANA ADVANCED RES & TECHNOLOGY.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     capsulatum rRNA ITS1 primer 1724F.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example 1; Col 5; 10pp; English.
                                                                                                                                                                                                                                                                                                                                       1297 AACGAGGAGTTCAAGAC 1313
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     622 AAGCTGGACAAACTGGG 638
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       BP.
                                                                                                                                                                                                                                                                                                                                                                                                AACCAGGAGCTCAAGAC 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1 AAGCTGGTCAAACTTGG 17
                                                                                                                                                                                                                      0.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0.8%;
                                                                                              prophylaxis of such diseases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         95US-00400580
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     95US-00400580
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAV01575 standard; DNA; 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
                                                                                                                                                                                                                                                                                  15; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Ajellomyces capsulatus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAX17891/c
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AAX17891 standard; DNA; 19

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replication. Optionally the oligomuclectides include at least one 2'-(2-methoxyethoxy) sugar modification or phosphorothicate internucleotide linkages. The oligonucleotides are used to inhibit CMV infections (by in vivo or in vitro contact with cells, tissues or body fluids), especially
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gibberellin 4; GA4; beta-hydroxylase; GA4 homologue; GA4H; GA4H1; GA4H2; plant growth hormone; seed germination; stem elongation; flowering; fruiting; stem growth; alpha-tubulin; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New antisense oligonucleotides that target cytomegalovirus nucleic acid particularly including 2-methoxyethoxy sugar modifications, especially for treating viral retinitis, with long-lasting retention in the retina.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Antisense oligonucleotides (AAX17861-X17924) are targeted to a nucleic acid (AAX17925-X17948) encoding IE (immediate early) 1 or 2, or DNA polymerase of cytomegalovirus (CMV) and are able to inhibit CMV
                                                                                                                        Antisense; oligonucleotide; immediate early; DNA polymerase; CMV; cytomegalovirus; inhibition; replication; sugar modification; phosphorothioate; infection; retinitis; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                to treat or prevent CMV infections, particularly retinitis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 19 BP; 0 A; 5 C; 4 G; 10 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                     Chapman S;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PCR primer Tua4R used to amplify alpha-tubulin.
                                                                                                                                                                                                                                                                                                                                                                                                    Anderson KP,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 7; Page 30; 99pp; English.
                                                                                            Anti-CMV oligonucleotide #5478.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             133 ATGAAGAAGATCAAACG 149
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           18 AAGAAGAAGAGCAAACG 2
                                                                                                                                                                                                                                                                                                       98WO-US006895
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                                                            (first entry)
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                                                                                                                                                                                                                                                                                                                                                                       (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 1998-568330/48.
                                                                                                                                                                                                           Human herpesvirus 5.
                                                                                                                                                                                                                                                                                                                                       09-APR-1997;
                                                                                                                                                                                                                                                                                                       07-APR-1998;
                                                                                                                                                                                                                                       WO9845314-A1
                                                            11-MAY-1999
                                                                                                                                                                                                                                                                       15-OCT-1998
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                                                                                                                                                                                        Synthetic
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                            AAX17891
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                                                                                                                                                                                                                                                                                                                             PCR primers AAX04626-27 were used to amplify the alpha-tubulin 4 gene. The primers are used as an internal control when determining expression of the GA4H1 gene. GA4H1 is a gibberellin 4 (GA4) homologue. The GA4H proteins (GA4H1 and GA4H2) have similar functions to GA4. GA4H is believed to be a member of the enzyme family involved in the biosynthesis of the gibberellin family of plant growth hormones that promote various growth and developmental processes in higher plants, such as seed germination, stem elongation, flowering and fruiting. GA4 is a betahydroxylase, and the homologues may also have 3-betahydroxylase activity, which is critical for controlling stem growth. GA4H may be applied to crops to enhance and facilitate stem elongation, flowering and fruiting. Alternatively, the DNA encoding GA4H may be genetically inserted into the plant host to produce a similar effect
                                                                                                                                                                                                                                    New isolated Gibberellin 4 homologues - derived from Arabidopsis plants, used to develop products for altering stem growth, e.g. for enhancing stem elongation, flowering and fruiting.
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88.2%; Pred. No. 9.8e+02;
ive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Probe hybridising to nucleotides of human c-erb-B-2 (HER-2)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human, c-erb-B-2; HER-2; chromosome aberration; probe;
peptide nucleic acid; haemapoietic malignancy; cancer;
inborn constituel disease; herbicide resistance gene; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 19 BP; 3 A; 6 C; 2 G; 8 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                      Example 5; Page 33; 106pp; English
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                                                                                                                                                                       Chiang
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1517 TAAAGGAGATTCAGCTA 1533
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                              98WO-US013044.
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                                                            97US-0050615P.
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                                                                                        (GEHO ) GEN HOSPITAL CORP. (GOOD/) GOODMAN H M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Best_Local Similarity 88.2
Matches 15; Conservative
                                                                                                                                                                        Goodman HM, Nguyen LV,
                                                                                                       (GOOD/) GOODMAN H M.
(NGUY/) NGUYEN L V.
(CHIA/) CHIANG H.
                                                                                                                                                                                                        WPI; 1999-105626/09
                              24-JUN-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    04-MAY-1999;
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                                                            24-JUN-1997;
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30-DEC-1998
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designed to cleave RNA encoding a cyclin or cell-cycle dependent kinase other than cell-cycle dependent kinases CDK1, PCNA and Cyclin B1. Representative examples of ribozyme recognition sites are given in AAA82415 to AAA86787. The ribozyme of the invention is useful for inhibiting restenosis by introduction of the ribozyme into cells. The ribozyme is resistant to endonuclease activity and hence is efficient in restenosis treatment
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The present invention relates to a hairpin or hammerhead ribozyme, designed to cleave RNA encoding a cyclin or cell-cycle dependent kinase other than cell-cycle dependent kinases CDK1, PCNA and Cyclin B1. Representative examples of ribozyme recognition sites are given in AAA82415 to AAA86787. The ribozyme of the invention is useful for inhibiting restenosis by introduction of the ribozyme into cells. The ribozyme is resistant to endonuclease activity and hence is efficient in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Ribozyme; hairpin; hammerhead; gene therapy; vasotropic; restenosis; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New hairpin and hammerhead ribozyme for inhibiting restenosis, cleaves RNA encoding a cyclin or cell-cycle dependent kinase other than CDK1,
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                                                                                                                                                                                                                            Length 19;
                                                                                                                                                                                                                                                                      2; Indels
                                                                                                                                                                                 Sequence 19 BP; 7 A; 6 C; 3 G; 3 T; 0 U; 0 Other;
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                                                                                                                                                                                                                          Score 13.8; DB 1;
Pred. No. 9.8e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Robbins JM;
                                                                                                                                                                                                                                                                   0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Barber JR,
                                                                                                                                                                                                                                                                                                               1138 TACTCCACTCAGATTGA 1154
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    cdk4 ribozyme binding site #55
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           973 CACCGAGACCTCAAGCC 989
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                                                                                                                                                                                                                          0.8%;
                                                                                                                                                                                                                                                                                                                                                          1 TACTCCACTCAGAAGA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAA82874 standard; DNA; 19
                                                                                                                                                                                                                                                                      15; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PCNA and Cyclin Bl.
                                                                                                                                                                                                                                               Local Similarity
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                                                                                                                                                                                                                   coligonucleotides AAZ36562-97 represent a set of probes hybridising to the human c-exb-B-2 (HER-2) gene. The probes are used to demonstrate the method of the invention. The specification describes a method for the detection of chromosome aberrations in eukaryotic samples uses sets of peptide nucleic acid (PNA) probes in hybridisation reactions. The method comprises using at least 2 sets of hybridisation reactions. The method composities using at least 2 sets of hybridisation probes, where at least composine. The method sequences related to a potential aberration in a chromosome. The methods can be used for the detection of chromosome aberrations. They can be used for the diagnosis of disorders and diseases related to chromosomal aberrations or abnormalities such as e.g. hampoietic malignancies, cancers and inborn constituel diseases. The method may be used for detecting viral sequences and their localization in the chromosome. In plant biology, the methods can be used for monitoring the efficiency of transferring herbicide resistance genes to a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0;
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                                                                                             of chromosome aberrations, used for detecting diseases and infections, and plant alterations related to e.g. herbicide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  cleaves
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The present invention relates to a hairpin or hammerhead ribozyme,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 19 BP; 3 A; 5 C; 5 G; 6 T; 0 U; 0 Other;
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           Adelhorst K;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Disclosure; Page 46; 109pp; English.
                                                                                                                                                                                   Example 1; Page 44; 63pp; English
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        cdk1 ribozyme binding site #20.
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         Nielsen KV,
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Matches 15; Conserv
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                                                                                             Detection of
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                                                                                                                                          resistance.
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           Pluzek K,
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The present invention relates to a hairpin or hammerhead ribozyme, designed to cleave RNA encoding a cyclin or cell-cycle dependent kinase other than cell-cycle dependent kinases CDK1, PCNA and Cyclin B1.
Representative examples of ribozyme recognition sites are given in AAA82415 to AAA86787. The ribozyme of the invention is useful for inhibiting restenosis by introduction of the ribozyme into cells. The ribozyme is resistant to endonuclease activity and hence is efficient in restenosis treatment
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                                                                                                                                                        New hairpin and hammerhead ribozyme for inhibiting restenosis, cleaves RNA encoding a cyclin or cell-cycle dependent kinase other than CDK1, PCNA and Cyclin B1.
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18.2%; Pred. No. 9.8e+02;
ve 0; Mismatches 2; Indels
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                                                                                          Robbins JM;
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                                                                                                                                                                                                                                           Disclosure; Page 76; 109pp; English.
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                                                                                          Barber JR,
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                                                     (IMMU-) IMMUSOL INC.
                                                                                          Welch PJ,
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               04-DEC-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The present invention relates to a hairpin or hammerhead ribozyme, designed to cleave RNA encoding a cyclin or cell-cycle dependent kinase other than cell-cycle dependent kinases CDK1, PCNA and Cyclin Bl.
Representative examples of ribozyme recognition sites are given in AAA82415 to AAA86787. The ribozyme of the invention is useful for inhibiting restenosis by introduction of the ribozyme into cells. The ribozyme is resistant to endonuclease activity and hence is efficient in restenosis treatment
                                                                                                                                                                                     Ribozyme; hairpin; hammerhead; gene therapy; vasotropic; restenosis; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                new hairpin and hammerhead ribozyme for inhibiting restenosis, cleaves RNA encoding a cyclin or cell-cycle dependent kinase other than CDKI, PCNA and Cyclin B1.
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88.2%; Pred. No. 9.8e+02;
iive 0; Mismatches 2; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                        Barber JR,
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                                                                                                                                                cdk3 ribozyme binding site #14
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                                   AAA82729 standard; DNA; 19
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Matches 15; Conserv
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Ribozyme; hairpin; hammerhead; gene therapy; vasotropic; restenosis; ss
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Representative examples of ribozyme recognition sites are given in AAA88415 to AAA86787. The ribozyme of the invention is useful for inhibiting restenosis by introduction of the ribozyme into cells. The ribozyme is resistant to endonuclease activity and hence is efficient in
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Pred. No. 9.8e+02;
0; Mismatches 2;
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                                                                                                                                                                                                                                                                                                                                                                             Barber JR,
                                                                                       cdk3 ribozyme binding site #33
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Best Local Similarity 88.2
Matches 15; Conservative
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                                                                                                                                                                                                                                                                                                                                                                             Tritz R, Welch PJ,
                                                                                                                                                                                                                                                                                                                                         (IMMU-) IMMUSOL INC
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                  AAA82748;
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                                     The present invention relates to a hairpin or hammerhead ribozyme, designed to cleave RNA encoding a cyclin or cell-cycle dependent kinase other than cell-cycle dependent kinases CDK1. PCNA and Cyclin B1. Representative examples of ribozyme recognition sites are given in AAA82415 to AAA86787. The ribozyme of the invention is useful for inhibiting restenosis by introduction of the ribozyme into cells. The ribozyme is resistant to endonuclease activity and hence is efficient in restenosis treatment
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    hairpin; hammerhead; gene therapy; vasotropic; restenosis; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New hairpin and hammerhead ribozyme for inhibiting restenosis, cleaves RNA encoding a cyclin or cell-cycle dependent kinase other than CDK1, PCNA and Cyclin B1.
AAA82415 to AAA86787. The ribozyme of the invention is useful for inhibiting restenosis by introduction of the ribozyme into cells. The ribozyme is resistant to endonuclease activity and hence is efficient restenosis treatment
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Pred. No. 9.8e+02;
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0.8%; Score 13.8; DB 1; Length 19;
Best Local Similarity 88.2%; Pred. No. 9.8e+02;
Matches 15; Conservative 0; Mismatches 2; Indels
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                                                                                        Sequence 19 BP; 4 A; 5 C; 4 G; 6 T; 0 U; 0 Other
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                                                                                                                                                                                                1090 GTGACACTGTGGTACCG 1106
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                                                                                                                                                                                                                                                                                                                                                                                                                               cdk6 ribozyme binding site #80.
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Best Local Similarity 88.2%;
Matches 15; Conservative
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restenosis treatment
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Best Local Similarity
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                                                                                                                                                                                                                             04-MAY-2001
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                                                                                                                                        The present invention relates to a hairpin or hammerhead ribozyme, designed to cleave RNA encoding a cyclin or cell-cycle dependent kinase other than cell-cycle dependent kinases CDK1, PCNA and Cyclin Bl. Representative examples of ribozyme recognition sites are given in AA82415 to AAA86787. The ribozyme of the invention is useful for inhibiting restenosis by introduction of the ribozyme into cells. The ribozyme is resistant to endonuclease activity and hence is efficient in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The present invention relates to a hairpin or hammerhead ribozyme, designed to cleave RNA encoding a cyclin or cell-cycle dependent kinase other than cell-cycle dependent kinases CDKI, PCNA and Cyclin B1. Representative examples of ribozyme recognition sites are given in AAABS415 to AAAB6787. The ribozyme of the invention is useful for inhibiting restenosis by introduction of the ribozyme into cells. The ribozyme is resistant to endonuclease activity and hence is efficient in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Ribozyme; hairpin; hammerhead; gene therapy; vasotropic; restenosis; ss.
                                                               New hairpin and hammerhead ribozyme for inhibiting restenosis, cleaves RNA encoding a cyclin or cell-cycle dependent kinase other than CDKI, PCNA and Cyclin B1.
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RNA encoding a cyclin or cell-cycle dependent kinase other than CDKL,
PCNA and Cyclin B1.
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                       Robbins JM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Robbins JM
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                                                                                                                   Disclosure; Page 49; 109pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Welch PJ, Barber JR,
                      Barber JR,
                                                                                                                                                                                                                                                                                                                         1022 TCAAGCTGGCTGACTTT 1038
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                                                                                                                                                                                                                                                                                                Conservative
                                                                                                                                                                                                                          restenosis treatment
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Best Local Similarity
Thes 15, Conserve
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                      Welch PJ,
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(IMMU-) IMMUSOL INC.
                                            WPI; 2000-412314/35
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                      Tritz R,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The present invention provides nucleotides encoding molecular variants of the human multi drug resistance-1 (MDR-1) protein. These can be used to identify compounds capable of treating multidrug resistance and sensitivity interfering resulting from polymorphisms in MDR-1, which can lead to difficulties in treating cancer, cardiovascular, neuronal, inflammatory and CNS diseases
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                                                                                                                   Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human, MDR-1; multi drug resistance-1; drug uptake; disease; cancer; inflammatory disease; neuronal disease; CNS disease; cardiovascular disease; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human multi drug resistance-1 gene related sequence SEQ ID NO: 289
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                                                   Score 13.8; DB 1; Length 19;
Pred. No. 9.8e+02;
0; Mismatches 2; Indels
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Sequence 19 BP; 2 A; 8 C; 3 G; 6 T; 0 U; 0 Other;
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                                                      0.8%;
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22-FEB-2000; 2000EP-00103361
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nes 15; Conservative
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                                                                                                                                                                                                                                                                                                                                                                     The present invention provides nucleotides encoding molecular variants of the human multi drug resistance-1 (MDR-1) protein. These can be used to identify compounds capable of treating multidrug resistance and sensitivity interfering resulting from polymorphisms in MDR-1, which can lead to difficulties in treating cancer, cardiovascular, neuronal, inflammatory and CNS diseases
                                                                                                                                                                                                                                                                                                    New polynucleotide encoding a molecular variant Multi Drug Resistance (MDR)-1 polypeptide is useful for diagnosing and treating diseases associated with abnormal MDR-1 expression or function, e.g. cancer.
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                                                         Human; MDR-1; multi drug resistance-1; drug uptake; disease; cancer; inflammatory disease; neuronal disease; CNS disease; cardiovascular disease; PCR primer; ss.
                                   Human multi drug resistance-1 gene related sequence SEQ ID NO: 291.
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                                                                                                                                                                                                                                                                                                                                                    Disclosure; Page 137; 154pp; English
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Best Local Similarity 78.9
Matches 15; Conservative
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                                                                                                       Homo sapiens
            04-MAY-2001
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                                                                                                                                                  08-FEB-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Synthetic
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The present invention describes a method for treating a proliferative skin or eye disease and scarring. The method involves administering a ribozyme [1] which cleaves RNA encoding a cytokine involved in inflammation, matrix metalloproteinase (MMP), cyclin, cell-cycle dependent kinase, growth factor or a reductase, or administering a nucleic acid modecule [11] comprising a promoter operably linked to a nucleic acid segment encoding [12]. [1] can have antipsoriatic, dermatological, cytostatic, antiseborrheic, antidiabetic, antisickling, cleaves RNA encoding cytokine involved in inflammation. [1] can be used in gene therapy. [1] and [11] are useful for treating proliferative skin diseases such as psoriasis, atopic dermatitis, actinic keratosis, squamous or basal cell carcinoma and viral or seborrheic wart. They can also be used for treating proliferative eye diseases such as diabetic retinopathy, vitreoretinopathy, sickle cell retinopathy, retinopathy of prematurity and retinal decachment, and for treating and preventing content scarring such as keloid, adhesion and hypertrophic or hypertrophic burn scar. AAHS7577 to AAH62099 represent sequences used in the
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                                                                                                                                                                                                                                                                                                                                                                                                                                                    Treating proliferative skin or eye diseases and scarring, using ribozymes that cleave RNA encoding cytokines involved in inflammation, matrix metalloproteinases, growth factors and cell-cycle dependent kinases.
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88.2%; Pred. No. 9.8e+02;
ve 0; Mismatches 2; Indels
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26-OCT-2000; 2000WO-US029500
                                                                                       99US-0161532P
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Les 15; Conservative
                                                                                                                                                                                                                                                                          Tritz R;
                                                                                                                                                                                                                                                                                                                                                                 WPI; 2001-300427/31.
                                                                                                                                                                                   (IMMU-) IMMUSOL INC.
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26-OCT-2000; 2000WO-US029500.
 sickle cell retinopathy; ss.
                                                                      WO200130362-A2
                              Homo sapiens.
                                                                                                    03-MAY-2001.
                                           Synthetic
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The present invention describes a method for treating a proliferative skin or eye disease and scarring. The method involves administering a ribozyme (I) which cleaves RNA encoding a cytokine involved in inflammation, matrix metalloproteinase (MMP). cyclin, cell-cycle dependent kinase, growth factor or a reductase, or administering a nucleic acid molecule (II) comprising a promoter operably linked to a nucleic acid molecule (II) comprising a promoter operably linked to a nucleic acid molecule (II) comprising a promoter operably linked to a nucleic acid molecule (II) comprising a promoter operably linked to a nucleic acid molecule (II) comprising a promoter operably linked to a nucleic acid molecule (II) ace useful for treating proliferative skin diseases such as psoriasis, atopic dermatitis, actinic keratosis, squamous or basal cell carcinoma and viral or seborrheic wart. They can also be used for treating proliferative eye diseases such as diabetic retinopathy, vitreoretinopathy, sickle cell retinopathy, retinopathy of prematurity and retinal detachment, and for treating and preventing scarring such as kelloid, adhesion and hypertrophic or hypertrophic burn
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                                                                                                                                                                                                                                                 Treating proliferative skin or eye diseases and scarring, using ribozymes that cleave RNA encoding cytokines involved in inflammation, matrix metalloproteinases, growth factors and cell-cycle dependent kinases.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
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88.2%; Pred. No. 9.8e+02;
ative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             scar. AAH57577 to AAH62099 represent sequences used in the
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                                                                                                    26-OCT-2000; 2000WO-US029500.
                                                                                                                               99US-0161532P
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                                                                                                                                                                                        Robbins JM, Tritz R;
                                                                                                                                                           (IMMU-) IMMUSOL INC.
                                                                                                                                                                                                                    WPI; 2001-300427/31.
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Best Local Similarity
Matches 15; Conserv
                                           WO200130362-A2.
Homo sapiens.
Synthetic.
                                                                                                                                 26-OCT-1999;
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The present invention describes a method for treating a proliferative skin or eye disease and scarring. The method involves administering a ribozyme [1] which cleaves RNA encoding a cytokine involved in inflammation, matrix metalloproteinase (MMP), cyclin, cell-cycle dependent kinase, growth factor or a reductase, or administering a nucleic acid molecule [11] comprising a promoter operably linked to a concleic acid segment encoding [1]. [1] can have antiposoriatic, dermatological, cytostatic, antiseborrheic, antidiabetic, antisickling, ophthalmological, vulnerary, keratolytic and virucide activities, and cleaves RNA encoding cytokine involved in inflammation. [1] can be used in gene therapy. [1] and [11] are useful for treating proliferative skin diseases such as psoriasis, atopic dermatitis, actinic keratosis, suppose the control of treating proliferative eye diseases such as diabetic retinopathy, vitreoretinogathy, sickle cell retinopathy, retinopathy of prematurity and retinal detechment, and for treating and preventing proliferative of the diseases such as diabetic prematurity and retinal detechment, and for treating and preventing proliferative of the diseases and activities.
                                                                                                                                                                                                                                                                                                                                                            Treating proliferative skin or eye diseases and scarring, using ribozymes that cleave RNA encoding cytokines involved in inflammation, matrix metalloproteinases, growth factors and cell-cycle dependent kinases.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             such as keloid, adhesion and hypertrophic
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99US-0161532P.
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                                                                                                                                                                              Robbins JM, Tritz R;
                                                                                         (IMMU-) IMMUSOL INC.
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26-OCT-1999;
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The present invention describes a method for treating a proliferative skin or eye disease and scarring. The method involves administering a ribozyme [1] which cleaves RNA encoding a cytokine involved in inflammation, matrix metalloproteinase (MMP), cyclin, cell-cycle dependent kinase, growth factor or a reductase, or administering a nucleic acid molecule [II] comprising a promote operably linked to a concleic acid segment encoding [1]. [1] can have antipsoriatic, dermatological, cytostatic, antiseborrheic, antidiabetic, antisickling, ophthalmological, vulnerary, keratolytic and virucide activities, and cleaves RNA encoding cytokine involved in inflammation. [1] can be used in gene therapy. [1] and [II] are useful for treating proliferative skin diseases such as psoriasis, atopic dermatitis, actinic keratosis, squamous or basal cell carcinoma and viral or seborrheic wart. They can also be used for treating proliferative eye diseases such as diabetic retinopathy, vitreoretinopathy, sickle cell retinopathy, retinopathy of prematurity and retinal detechment, and for treating and preventing.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Treating proliferative skin or eye diseases and scarring, using ribozymes that cleave RNA encoding cytokines involved in inflammation, matrix metalloproteinases, growth factors and cell-cycle dependent kinases.
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recognition site, target, ribozyme binding site, eye disease, vulnerary,
proliferative disease, skin disease, psoriasis, diabetic retinopathy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        scarring such as keloid, adhesion and hypertrophic or hypertrophic burn scar. AAH57577 to AAH62099 represent sequences used in the
antipsoriatic, dermatological, antiseborrheic, antidiabetic, virucide, antisickling, ophthalmological, keratolytic, gene therapy, viral wart, atopic dermatitis, actinic keratosis, squamous cell carcinoma,
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                                                                                                  basal cell carcinoma; seborrheic wart; vitreoretinopathy; scar; sickle cell retinopathy; ss.
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Conservative
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                                                                                                                                                                                                                                                                                                        WO200130362-A2.
                                                                                                                                                                                                            Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   26-OCT-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Robbins JM,
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                                                                                                                                                                                                                                          Synthetic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Treating proliferative skin or eye diseases and scarring, using ribozymes that cleave RNA encoding cytokines involved in inflammation, matrix metalloproteinases, growth factors and cell-cycle dependent kinases.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       scarring such as Keloid, adhesion and hypertrophic or hypertrophic burn scar. AAHS7577 to AAHS2099 represent sequences used in the
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      atopic dermatitis, actinic keratosis, squamous cell carcinoma; basal cell carcinoma; seborrheic wart; vitreoretinopathy; scar; sickle cell retinopathy; ss.
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88.2%; Pred. No. 9.8e+02;
ve 0; Mismatches 2; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           exemplification of the present invention
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Best Local Similarity 88.2%;
Matches 15; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Tritz R;
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                                                                                                                                                                                                                                       WO200130362-A2
                                                                                                                                              sapiens.
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                                                                                                                                                                          Synthetic
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recognition site; target; ribozyme binding site; eye disease; vulnerary; proliferative disease; skin disease; psoriasis; diabetic retinopathy; cytokine; inflammation; cell-cycle dependent kinase; cyclin; MMP; matrix metalloproteinase; growth factor; reductase; scarring; cytostatic; antipsoriatic; dermatological; antiseborrheic; antidiabetic; virucide; antisicking; ophthalmological; keratolytic; gene therapy; viral wart; atopic dermatitis; actinic keratosis; squamous cell carcinoma; basal cell carcinoma; seborrheic wart; vitreoretinopathy; scar; sickle cell retinopathy; ss.

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The present invention describes a method for treating a proliferative skin or eye disease and scarring. The method involves administering a ribozyme (1) which cleaves RNA encoding a cytokine involved in inflammation, matrix metalloproteinase (MPD), cyclin, cell-cycle dependent kinase, growth factor or a reductase, or administering a nucleic acid molecule (II) comprising a promoter operably linked to a cucleic acid segment encoding (1). (1) can have antipsoriatic, anticlaseborrheic, anticlabetic, anticlabetic, anticlabetic, anticlabetic, anticlamentological, cytostatic, antiseborrheic, anticlamenton. (1) can be used in gene therapy. (1) and (II) are useful for treating proliferative skin diseases such as psoriasis, atopic dermatitis, actinic keratosis, squamous or basal cell carcinoma and viral or seborrheic wart. They can also be used for treating proliferative eye diseases such as diabetic retinopathy, vitreoretinopathy, sickle cell retinopathy, retinopathy of prematurity and retinal detachment, and for treating and preventing
cytokine; inflammation; cell-cycle dependent kinase; cyclin; MMP; matrix meteralloproteinase; growth factor; reductase; scarring; cytostatic; antipocial; classe; antipocial; antiseborrheic; antidiabetic; virucide; antisickling; ophthalmological; keratolytic; gene therapy; viral wart; actopic dermatitis, actinic keratosis; squamous cell carcinoma; basal cell carcinoma; seborrheic wart; vitreoretinopathy; scar; sickle cell retinopathy; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Treating proliferative skin or eye diseases and scarring, using ribozymes that cleave RNA encoding cytokines involved in inflammation, matrix metalloproteinases, growth factors and cell-cycle dependent kinases.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              scarring such as keloid, adhesion and hypertrophic or hypertrophic scar. AAH57577 to AAH62099 represent sequences used in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          exemplification of the present invention
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                                                                                                                                                                                                                                                                                                                                                                                    99US-0161532P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2001-300427/31.
                                                                                                                                                                                                                                                                                                                                                                                                                                (IMMU-) IMMUSOL INC.
                                                                                                                                                                                                                                                  WO200130362-A2.
                                                                                                                                                                                   sapiens.
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                                                                                                                                                                                                                                                                                              33-MAY-2001
                                                                                                                                                                                                        Synthetic.
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Treating proliferative skin or eye diseases and scarring, using ribozymes that cleave RNA encoding cytokines involved in inflammation, matrix metalloproteinases, growth factors and cell-cycle dependent kinases.

26-OCT-2000; 2000WO-US029500.

WO200130362-A2.

03-MAY-2001.

sapiens.

Homo

Synthetic.

99US-0161532P.

26-OCT-1999;

Tritz R;

Robbins JM,

(IMMU-) IMMUSOL INC.

WPI; 2001-300427/31.

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Kin or eye disease and scarring. The method involves administering a ribozyme (I) which cleaves RNA encoding a cytokine involved in inflammation, matrix metalloproteinase (MMP), cyclin, cell-cycle dependent kinase, growth factor or a reductase, or administering a nucleic acid molecule (II) comprising a promoter operably linked to a nucleic acid molecule (II) comprising a promoter operably linked to a nucleic acid molecule (II) comprising a promoter operably linked to a nucleic acid molecule (II) comprising a promoter operably linked to a nucleic acid molecule (II) comprising a promoter operably linked to a nucleic acid molecule (II) comprising a nutleice activities, and clematological, vulnerary, keratolytic and viruide activities, and cleaves RNA encoding cytokine involved in inflammation. (I) can be used in gene therapy. (I) and (II) are useful for treating proliferative skin diseases such as psoriasis, atopic dermatitis, actinic keratosis, squamous or basal cell carlinoma and viral or sebortheic wart. They can also be used for treating proliferative eye diseases such as diabetic retinopathy, vitreoretinopathy, sickle cell retinopathy, retinopathy of prematurity and retinal detechment, and for treating and preventing scarring such as keloid, adhesion and hypertrophic or hypertrophic burn scar. AAH57577 to AAH62099 represent sequences used in the cemplification of the present invention
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0
                             0.8%; Score 13.8; DB 1; Length 19;
                                         Pred. No. 9.8e+02;
0; Mismatches 2; Indels
Sequence 19 BP; 4 A; 5 C; 4 G; 6 T; 0 U; 0 Other;
                                                                                      1090 GTGACACTGTGGTACCG 1106
                                                                                                                 18
                                                                                                                                                                                        BP
                                         88.2%;
                                                                                                                 2 GTTACACTCTGGTACCG
                                                                                                                                                                                       AAH57596 standard; DNA; 19
                                                                                                                                                                                                                                                (first entry)
                                           Best Local Similarity 88.2
Matches 15; Conservative
                                                                                                                                                                                                                                                10-SEP-2001
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                             Query Match
                                                                                                                                                            RESULT 1426
                                                                                                                                                                          AAH57596
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Cell-cycle dependent kinase cdk4 ribozyme binding site SEQ ID NO:473.

10-SEP-2001 (first entry)

AAH58049;

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AAH58049
ID AAH5
XX
AC AAH5
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DT 10-S
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Human; ribozyme therapy; hairpin ribozyme; hammerhead ribozyme;

matrix metalloproteinase; growth factor; reductase; scarring; cytostatic; antipsoriatic; dermatological; antiseborrheic; antidiabetic; virucide; attisic, copthalmological; keratolytic; gene therapy; viral wart; atopic dermatitis; actinic keratosis; squamous cell carcinoma; basal cell carcinoma; seborrheic wart; vitreoretinopathy; scar; sickle cell retinopathy; ss.

26-OCT-2000; 2000WO-US029500.

WO200130362-A2. Homo sapiens. Synthetic.

03-MAY-2001,

99US-0161532P.

26-OCT-1999;

Robbins JM, Tritz R; (IMMU-) IMMUSOL INC.

Human, ribozyme therapy; hairpin ribozyme; hammerhead ribozyme; recognition site; target; ribozyme binding site; eye disease; vulnerary; proliferative disease; skin disease; psoriasis; diabetic retinopathy; cytokine; inflammation; cell-cycle dependent kinase; cyclin, MMP;

Cell-cycle dependent kinase cdk3 ribozyme binding site SEQ ID NO:335.

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The present invention describes a method for treating a proliferative skin or eye disease and scarring. The method involves administering a cribozyme (I) which cleaves RNA encoding a cytckine involved in inflammation, matrix metalloproteinase (MMP), cyclin, cell-cycle dependent kinase, growth factor or a reductase, or administering a nucleic acid molecule (II) comprising a promoter operably linked to a culcie acid segment encoding (I). (I) can have antipsoriatic, antisickling, ophthalmological, cytostatic, antiseborrheic, antidiabetic, antisickling, ophthalmological, vulnerary, keratolytic and virucide activities, and cleaves RNA encoding cytokine involved in inflammation. (I) can be used in gene therapy. (I) and (II) are useful for treating proliferative skin diseases such as psoriasis, atopic dermatitis, actinic keratosis, squamous or basal cell carcinoma and viral or seborrheic wart. They can also be used for treating proliferative eye diseases such as diabetic retinopathy, vitreoretinopathy, sickle cell retinopathy, retinopathy of prematurity and retinal detachment, and for treating and preventing
              Human, ribozyme therapy; hairpin ribozyme; hammerhead ribozyme; recognition site; target; ribozyme binding site; eye disease; vulnerary; proliferative disease; skin disease; psoriasis; diabetic retinopathy; cytokine; inflammation; cell-cycle dependent kinase; cyclin; MMP; matrix metalloproteinase; growth factor; reductase; scarring; cytostatic; antipsoriatic; dermatological; antiseborrheic; antidiabetic; virucide; antisicking; ophthalmological; keratolytic; gene therapy; viral wart; atopic dermatitis; actinic keratosis; squamous cell carcinoma; basal cell carcinoma;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Treating proliferative skin or eye diseases and scarring, using ribozymes that cleave RNA encoding cytokines involved in inflammation, matrix metalloproteinases, growth factors and cell-cycle dependent kinases.
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                                                                                                                                                                                                                    sickle cell retinopathy; ss.
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                                                                                                                                                                                                                                                                     Homo sapiens
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                                                                                                                                                                                                                                                                                          Synthetic.
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0; scarring such as keloid, adhesion and hypertrophic or hypertrophic burn scar. AAH57577 to AAH62099 represent sequences used in the exemplification of the present invention 0.8%; Score 13.8; DB 1; Length 19; 88.2%; Pred. No. 9.8e+02; Sequence 19 BP; 7 A; 6 C; 3 G; 3 T; 0 U; 0 Other; 88.28;

Gaps ; Indels Mismatches ; 1138 TACTCCACTCAGATTGA 1154 Local Similarity 88.2 hes 15; Conservative Query Match Matches ò

TACTCCACTCAGAAAGA 17

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AAH57911 standard; DNA; 19 BP. (first entry) 10-SEP-2001 AAH57911;

1427

AAH57911

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The present invention describes a method for treating a proliferative skin or eye disease and scarring. The method involves administering a ribozyme (1) which cleaves RNA encoding a cytokine involved in inflammation, matrix metalloproteinase (WMP), cyclin, cell-cycle dependent kinase, growth factor or a reductase, or administering a nucleic acid molecule (II) comprising a promoter operably linked to a concleic acid molecule (II) comprising a promoter operably linked to a cucleic acid molecule (II) comprising a promoter operably linked to a cucleic acid segment encoding (I). (I) can have antipsoriatic, dermatological, cytostatic, antiseborrheic, antidiabetic, antisickling, ophthalmological, vulnerary, keratolytic and virucide activities, and cleaves RNA encoding cytokine involved in inflammation. (I) can be used in gene therapy. (I) and (II) are useful for treating proliferative skin diseases such as psoriasis, atopic dermatitis, actinic keratosis, squamous or basal cell carcinoma and viral or seborrheic wart. They can also be used for treating proliferative eye diseases such as diabetic retinopathy, vitreoretinopathy, sickle cell retinopathy, retinopathy of prematurity and retinial detechment, and for treating and preventing proverting the prematurity and retinial detechment, and for treating and preventing proverting and preventing proverting and preventing and preventing proverting and preventing proverting and preventing proverting and preventing and preventing proverting and preventing preventing and preventing and prevent
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                                                                                                        Treating proliferative skin or eye diseases and scarring, using ribozymes that cleave RNA encoding cytokines involved in inflammation, matrix metalloproteinases, growth factors and cell-cycle dependent kinases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    scarring such as keloid, adhesion and hypertrophic or hypertrophic burn scar. AAH57577 to AAH62099 represent sequences used in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ·.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 13.8; DB 1;
Pred. No. 9.8e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        exemplification of the present invention
                                                                                                                                                                                                                                                                                                                                    Example 1; Page 96; 408pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    935
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TACCICITCCAGCIGCT 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABS67829 standard; DNA; 19 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0.8%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            15; Conservative
WPI; 2001-300427/31.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity
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AC ABS6
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The present invention relates to antisense oligonuclectides and methods for modulating the expression of human or mouse casein kinase 2-alpha prime. The antisense oligonuclectides are useful for inhibiting the expression of casein kinase 2-alpha prime, and for treating diseases or conditions associated with aberrant expression of casein kinase 2-alpha prime, and for treating diseases or prime. Such diseases include diabetes mellitus, and hyperproliferative disorders (particularly cancers e.g. breast cancer, prostate cancer, or liver cancer). The antisense compounds are also useful for diagnostics, therapeutics, prophylaxis, e.g. to prevent or delay infection, inflammation or tumour formation, as research reagents and kits, and in distinguishing between functions of various members of a biological pathway. The present sequence represents a PCR primer used to amplify DNA encoding human casein kinase 2-alpha prime in the examples of the present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Chinese hamster; expression augmenting sequence element; EASE; HMG-I(Y); recombinant protein expression; mammalian host cell; PCR; primer; ss; high mobility group; nonhistone chromatin protein; architectural transcription factor.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New antisense oligonucleotides targeted to nucleic acid encoding casein kinase 2-alpha prime, useful for diagnosing and/or treating a disease or condition associated with expression of casein kinase 2-alpha prime.
                                                                                    Human, casein kinase 2-alpha prime, diabetes mellitus;
hyperproliferative disorder, breast cancer, prostate cancer,
liver cancer, infection, inflammation, tumour formation, cytostatic,
antidiabetic, antiinflammatory, antimicrobial, PCR, primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
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0
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Pred. No. 9.8e+02;
0; Mismatches 2; Indels
                                                      Human casein kinase 2-alpha prime DNA, PCR primer #2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 19 BP; 1 A; 8 C; 3 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Chinese hamster HMG-I(Y) PCR primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example 13; Page 91; 129pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1364 GACTTGATAGCGACGGG 1380
                                                                                                                                                                                                                                                                                                                                                                                                                            Wyatt JR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAK98357 standard; DNA; 19 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0.8%;
88.2%;
                                                                                                                                                                                                                                                                                                           01-FEB-2002; 2002WO-US002772.
                                                                                                                                                                                                                                                                                                                                               08-FEB-2001; 2001US-00780173
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GACTGGAAAGCGACGGG 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity
Local Sconservative
Local 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  08-MAY-2002 (first entry)
              29-NOV-2002 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                     (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                            Freier SM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2002-627539/67.
                                                                                                                                                                                                                              WO200262951-A2.
                                                                                                                                                                                           Homo sapiens
                                                                                                                                                                                                                                                                    15-AUG-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAK98357;
                                                                                                                                                                                                                                                                                                                                                                                                                            Mckay R,
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sequences of the invention contain numerous binding sites for members of the HMG-I(Y) ("high mobility group") family of nonhistone chromatin proteins, a group of minor groove-binding architectural transcription factors which are thought to be involved in the mechanisms by which EASE sequences improve expression of transgenes. The EASEs of the invention of transgenes. The EASEs of the invention of transfer and also be used in the identification of additional EASE sequences (e.g. from other transformed cell lines which exhibit high levels of expression not attributable to a high gene copy number). Expression of recombinant in microbial (prokaryotic) cells, since the post-translational modifications found in mammalian cells are more likely to resemble those
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   found in a mammal. The present sequence represents a Chinese hamster high mobility group nonhistone chromatin protein-I(Y) (HMG-I(Y)) PCR primer, used in an example of the invention to clone the Chinese hamster HMG-I(Y)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human; chromosome 1p36-35; chromosome 21q22.1; genetic analysis; genome;
                                                                                                                                                                                                                                                                                                                                                                                                           The invention comprises Chinese hamster expression augmenting sequence elements (EASEs; AAK98343-AAK98344) that can be used to improve expression of recombinant proteins in mammalian host cells. The EASE
                                                                                                                                                                                                                                                                                                       New expression augmenting sequence elements isolated from a Chinese hamster ovary cell line improve expression of recombinant proteins in host mammalian cells.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0.8%; Score 13.8; DB 1; Length 19;
88.2%; Pred. No. 9.8e+02;
ve 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human chromosome 1p36-35 PCR primer SEQ ID NO:744.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 19 BP; 2 A; 9 C; 3 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                           Example 16; Col 22; 25pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                406
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                                                                                                                                                     97US-00785150.
                                                                                                                                                                   99US-00435377.
                                                                                                   2000US-00660299.
                                                                                                                                   96US-00586509
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              88.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              390 CTCGGATGAGGTGCAGT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                19 CICGGAGGAGGAGCAGI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ABL43700 standard; DNA; 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
                                                                                                                                                                                                                                       Morris AE, Thomas JN;
                                                                                                                                                                                                     (IMMV ) IMMUNEX CORP.
                                                                                                                                                                                                                                                                         WPI; 2002-033281/04.
Cricetulus griseus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Local Similarity
hes 15; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             JP2001321190-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PCR primer; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens.
                                                                                                   12-SEP-2000;
                                 US6309841-B1
                                                                                                                                     11-JAN-1996;
                                                                                                                                                                     05-NOV-1999;
                                                                                                                                                       13-JAN-1997:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      11-APR-2002
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28-NOV-2001; 2001WO-US044838 28-NOV-2000; 2000US-00724389

WO200257410-A2 Homo sapiens.

25-JUL-2002

(DNAS-) DNA SCI LAB INC.

WPI; 2002-698522/75. Guida M, Hall J;

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The present invention describes a method of arraying genome clones. The method comprises: (a) clones of the genomic libraries contained in multiwell plates numbered for discrimination are mixed in each of the multiwell plates; (b) a primer designed based on the chromosome marker sequence is added to the mixture to carry out an amplification reaction; (c) a signal corresponding to the marker is detected from the resultant camplified product to specify the discrimination Nos. of the multiwell plates containing the clones having said marker sequence; (d) the order of the markers is changed so that the same discrimination Nos. succeed to the maximum in the specified discrimination Nos. to array the multiwell plates; (e) the clones in the multiwell plates of the specified discrimination Nos. to array the multiwell and lateral directions; (f) the mixed clones are cultured and the creat data from the amplified by using the above primer; (g) signals and lateral directions; (f) the mixed clones are cultured as the positions on the chromosome and arrayed. The microarray is useful for gene analysis. ABL42957 to ABL45322 represent PCR primers for human chromosome lages. ABL45322 represent PCR primers for human chromosome 21q22.1, which are specifically claimed for use in the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 19 BP; 5 A; 3 C; 9 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                              Claim 4; Page 19; 528pp; Japanese.
12-MAR-2001; 2001JP-00068285
                                                 10-MAR-2000; 2000JP-00066716
                                                                                             (RIKA ) RIKAGAKU KENKYUSHO.
(GENO-) GENOTEX YG.
                                                                                                                                                                                                                    Arraying genome clones.
                                                                                                                                                                     WPI; 2002-144136/19
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Gaps 0 0.8%; Score 13.8; DB 1; Length 19; 88.2%; Pred. No. 9.8e+02; lve 0; Mismatches 2; Indels 874 CTGGATGACTGTGGGAA 890 88.2%; Local Similarity 88.2 nes 15; Conservative Query Match Best Loca Matches ò

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ABS97865 standard; DNA; 19 BP (first entry) 23-DEC-2002 ABS97865; ABS97865/

Human UDP-glucuronosyl transferase 24B gene PCR primer #2.

Human, ss; primer; cytochrome P450 A1; CYP4501A1; UGT2B4; MDR1; PCR;

Cytochrome P450 A2; CYP4501A2; cytochrome P450 02E; CYP45002E1; LTF;

Adrenorgic receptor beta1; ADBR1; aryl hydrocarbon; AHR; MRS1; NRI12;

W adrenorgic receptor nuclear translocator; ARNT; cathepsin S; CTSS;

Cyclooxgenase 2; COX2; diazepam binding inhibitor; DBI; haematological;

W epoxide hydroxylase 2; EPHX2; 5-lipoxygenase activating protein; FLAP;

Glutathione-S-transferase 12; GST12; histamine-N-methyl transferase;

W MADPH quinone oxidoreductase 2; NQO2; sulfotransferase; NNMT;

W DP-glucuronosyl transferase 28; UDP-glucuronosyl transferase 2B7;

W UDP-glucuronosyl transferase; UGT2B15; usokinase receptor;

W UDP-glucuronosyl transferase; UGT2B15; usokinase receptor;

W ultidrug resistance 1; lactotransferrin; orphan nuclear receptor;

acetylcholine muscarinic receptor; CHMR1; CHMR2; CHMR3;

altered drug metabolism; cardiovascular function; colorectal tumour; central nervous system; pulmonary; immunological

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This invention relates to the sequence of an isolated nucleic acid molecule comprising at least one base variation from that of a known human cytochrome P450 Al (CYP4501A1), cytochrome P450 Al (CYP4501A1), cytochrome P450 O2E1 (CYP4501A1), advenergic receptor betal (ADBR1), cothergoin (ARN), cathegoin S (CTSS), cyclooxgenase 2 (CRX2), diazepam binding CC (ARNT), cathegoin S (CTSS), cyclooxgenase 2 (CRX2), diazepam binding cothergoin (ELAP), glutathione-S-transferase 1 (GYT12), histamine-N-methyl transferase (HNWT), NADPH quinone-Studioricase 2 (MOZ2), histamine-N-methyl cransferase (HNWT), NADPH quinone-Studioricase 2 (MOZ2), culfortansferase (HNWT), NADPH quinone-Studioricase 2 (MOZ2), culfortansferase (HNWT), nationase receptor (MRT2), or acetylofoline mancariation of cothergoin corporation and categorical propaga markers for locating and characterising the genetic linkage markers for locating and characterising the genetic linkage markers for locating and characterising the genes trasponsible for specific traits within the genome and eventually care responsible for specific traits within the genome and eventually care responsible for specific traits within the genome and eventually care responsible for specific acid molecules comprising the disorders. The nucleic acid molecules comprising the disorders. The nucleic acid molecules comprising and/or traits within the genome and eventually collymorphic sequences contained in CYP4501A1, CYPP4501A2, CYP4501A2, CYPA4501A1, CYPP4501A2, CYPA4501A2, CYPA4501A1, CYPA4501A2, CYPA4501A2, CYPA4501A2, CYPA4501A2, CYPA4501A3, CYPA4501A3, CYPA4501A3, CYPA4501A3, CYPA4501A3, CYPA4501A1, CYPA4501A3, CYPA4501
                                                                                                                                                                                                                                                                                                                                                                       Isolated nucleic acid molecules having polymorphisms in known human genes e.g. cytochrome p450 and cathepsin S useful as genetic linkage markers for locating, identifying and characterizing the genes responsible for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     present sequence represents a PCR
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 18; Page 133; 714pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                              disorder-related traits.
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0.8%; Score 13.8; DB 1; Length 19; 18.2%; Pred. No. 9.8e+02; ve 0; Mismatches 2; Indels 859 GACCTGAAGCAGTACCT 875 88.28; Query Match 0.8° Best Local Similarity 88.2 Matches 15; Conservative

Sequence 19 BP; 3 A; 4 C; 5 G; 7 T; 0 U; 0 Other;

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Gaps

0;

19 GACCTGAAGGAATACCT

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ABL95969;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The present invention relates to nucleic acid probes, which are useful for assaying nucleic acids by hybridising with a target nucleic acid, in which a single-stranded oligonucleotide is labelled with a fluorescent substance and a quencher in a manner that the fluorescence intensity of the hybridisation reaction system is increased after completion of the hybridisation but no stem loop structure is formed. The probes are useful for assaying nucleic acids and their polymorphism and mutation, particularly useful for e.g. analytical applications, disease diagnosis and microbial identification. The present sequence was used to illustrate
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                                                                                                                                                                           polymorphism detection; mutation detection; disease diagnosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Fluorescently-labeled nucleic acid probes for assaying nucleic acids their polymorphism and mutation, particularly useful in science and medicine for e.g. analytical applications, disease diagnosis and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Yamada
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0.8%; Score 13.8; DB 1; Length 19;
88.2%; Pred. No. 9.8e+02;
.ive 0; Mismatches 2; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 19 BP; 5 A; 5 C; 7 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (NAAD-) NAT INST ADVANCED IND SCI & TECHNOLOGY
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Probe #31 for assaying nucleic acids.
                                                                                                                                         Probe #46 for assaying nucleic acids
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ABL35971/C

ABL95971 standard; DNA; 19 BP
AC
ABL95971;
XX
ABL95971;
XX
19-JUN-2002 (first entry)
XX
Probe; polymorphism detection
XX
W
MC100208414-A1.
XX
WC200208414-A1.
XX
Sequence 19 BP; 5 A; 5 C; 7 C C
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03-AUG-2000; 2000JP-00236115.
26-SEP-2000; 2000JP-00292483.
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                                                                                                                                                                                                 microbial identification; ss
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nes 15; Conser
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substance and a quencher in a manner that the fluorescence intensity of the hybridisation reaction system is increased after completion of the hybridisation but no stem loop structure is formed. The probes are useful for assaying nucleic acids and their polymorphism and mutation, particularly useful for e.g. analytical applications, disease diagnosis and microbial identification. The present sequence was used to illustrate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The present invention relates to nucleic acid probes, which are useful for assaying nucleic acids by hybridising with a target nucleic acid, in which a single-stranded oligonucleotide is labelled with a fluorescent
                                                                                                                                                                                                                                                                                                                                                                   Yamada K;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Probe, polymorphism detection, mutation detection; disease diagnosis; microbial identification; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Fluorescently-labeled mucleic acid probes for assaying nucleic acids their polymorphism and mutation, particularly useful in science and medicine for e.g. analytical applications, disease diagnosis and microbial identification
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                                                                                                                                                                                                                                                                                                                                                                 Kanagawa T, Kamagata Y, Torimura M, Kurata S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0.8%; Score 13.8; DB 1; Length 19; 88.2%; Pred. No. 9.8e+02; ive 0; Mismatches 2; Indels
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03-AUG-2000; 2000JP-00236115.
26-SEP-2000; 2000JP-00292483.
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03-AUG-2000; 2000JP-00236115.
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Matches 15; Conservative
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                                               WO200208414-A1
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Unidentified.
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                                                                                                                                                                                                                                                                                                                                                                                       Yokomaku T;
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The present invention relates to nucleic acid probes, which are useful for assaying nucleic acids by hybridising with a target nucleic acid, in which a single-stranded oligonucleotide is labelled with a fluorescent substance and a quencher in a manner that the fluorescence intensity of the hybridisation reaction system is increased after completion of the hybridisation but no stem loop structure is formed. The probes are useful for assaying nucleic acids and their polymorphism and mutation, particularly useful for e.g. analytical applications, disease diagnosis and microbial identification. The present sequence was used to illustrate the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The present invention describes the use of irinotecan (I) or its derivative for the preparation of a pharmaceutical composition for treating colorectal, cervical, gastric, lung, ovarian or pancreatic cancer, or malignant glioma in a subject having a genome with a variant allele which comprises a cytochrome p450, subfamily IIIA (nifedipine cytostatic applications of (II) and (II) have cytostatic activity. The therapeutic applications of (I) is improved, since it is possible to individually treat a subject with an appropriate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New use of irinotecan for preparation of compositions for treating in subject having genome with variant allele comprising cytochrome subfamily IIIA, polypeptide 5 polynucleotide, termed CYP3A5.
their polymorphism and mutation, particularly useful in science amedicine for e.g. analytical applications, disease diagnosis and microbial identification.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Cancer based on CYP3A5 related oligonucleotide SEQ ID NO:471
                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 19;
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                                                                                                                                                                                                                                                                                                                                                                                                              0.8%; Score 13.8; DB 1;
88.2%; Pred. No. 9.8e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0; Mismatches
                                                                                                41; Page 103; 152pp; Japanese.
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2002EP-00011710.
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                                                                                                                                                                                                                              Fluorescently-labeled nucleic acid probes for assaying nucleic acids and their polymorphism and mutation, particularly useful in science and medicine for e.g. analytical applications, disease diagnosis and microbial identification.
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                                                                                                                     Yamada
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                                                                                                                     Kurata S,
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88.2%; Pred. No. 9.8e+02;
iive 0; Mismatches 2; Indels
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                                                                                                                     Torimura M,
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                                                 (NAAD-) NAT INST ADVANCED IND SCI & TECHNOLOGY.
(KANK-) KANKYO ENG CO LTD.
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(KANK-) KANKYO ENG CO LTD.
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                                                                                                                     Kamagata Y,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Kamagata Y,
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03-AUG-2000; 2000JP-00236115.
26-SEP-2000; 2000JP-00292483.
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         26-SEP-2000; 2000JP-00292483
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                                                                                                                     Kanagawa T,
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ABL95961

RESULT 1435

ABL95961 ID ABL

Query Match

Best Loc Matches

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cancer p450,

Yokomaku

Kurane

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Gaps

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Length 19; 3; Indels

Score 13.8; DB 1; Pred. No. 9.8e+02; 1; Mismatches

0.8%;

Query Match
Best Local Similarity 78.9
Matches 15; Conservative

388 TCCTCGGATGAGGTGCAGT 406

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TCCTCTGAGRATGTGCAGT

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The present invention describes the use of irinotecan (I) or its derivative for the preparation of a pharmaceutical composition for treating colorectal, gastric, lung, ovarian or pancreatic cancer, or malignant gliona in a subject having a genome with a variant alled which comprises a cytochrome p450, subfamily IIIA (nifedipine oxidase), polypeptide 5 (CYP3A5) polymucleotide (II). (I) and (II) have cytostatic activity. The therapeutic applications of (I) is improved, since it is possible to individually treat a subject with an appropriate derivative of (I). Therefore, undesirable, harmful or toxic effects are efficiently avoided. Unnecessary and potentially harmful treatment of those subjects who do not respond to the treatment with substances (nonresponders), as well as the development of drug resistances due to suboptimal drug dosing can be avoided. ACF62200 to ACF62751 and ABM34912 to ABM35013 represent sequences used in the
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dosage and/or an appropriate derivative of (I). Therefore, undesirable, harmful or toxic effects are efficiently avoided. Unnecessary and potentially harmful treatment of those subjects who do not respond to the treatment with substances (nonresponders), as well as the development of drug resistances due to suboptimal drug dosing can be avoided. ACF62200 to ACF62751 and ABM34912 to ABM35013 represent sequences used in the exemplification of the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New use of irinotecan for preparation of compositions for treating cancer in subject having genome with variant allele comprising cytochrome p450, subfamily IIIA, polypeptide 5 polynucleotide, termed CYP3A5.
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                                                                                                                                                                                                   0.8%; Score 13.8; DB 1; Length 19; 78.9%; Pred. No. 9.8e+02; tive 1; Mismatches 3; Indels
                                                                                                                                                              Seguence 19 BP; 3 A; 4 C; 5 G; 6 T; 0 U; 1 Other;
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24-MAY-2002; 2002EP-00011710.
                                                                                                                                                                                                                                                                                 388 TCCTCGGATGAGGTGCAGT
                                                                                                                                                                                                                                                                                                                                                                                                                         ACF62643 standard; DNA; 19
                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2003-268144/26.
                                                                                                                                                                                                                        Similarity
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                                                                                                                                                                                                 Query Match
Best Local Simi
Matches 15;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                              ACF62643;
                                                                                                                                                                                                                                                                                                                                                                                  RESULT 1437
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irinotecan; colorectal cancer; cervical cancer; gastric cancer; lung cancer; ovarian cancer; pancreatic cancer; malignant glioma; variant allele; multidrug resistance protein 1; MRP1; cytostatic; gene;

MRP1 based cancer related nucleic acid SEQ ID NO:471.

(first entry)

20-NOV-2003

ВР

ADB21313 standard; DNA; 19

ADB21313

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         present sequence represents
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              treating colorectal, cervical, gastric, lung, ovarian or pancreatic cancer, or malignant glioma in a subject, where the subject is a human (preferably African or Asian) or a mouse. The present sequence represents a sequence which is used in the exemplification of the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The present invention describes a method for the use of irinotecan (I) cits derivative for the preparation of a pharmaceutical composition for treating colorectal, cervical, gastric, lung, ovarian or pancreatic cancer, or malignant glioma in a subject having a genome with a variant allele which comprises a multidrug resistance protein 1 (MRP1) polynucleotide (II). (I) has cytostatic activity. (I) or its derivative can be used for the preparation of a pharmaceutical composition for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 19 BP; 3 A; 4 C; 5 G; 6 T; 0 U; 1 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  388 TCCTCGGATGAGGTGCAGT 406
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Use of irinotecan or its derivative for preparation of a pharmaceutical composition for treating cancer in a subject having a genome with a variant allele comprising a multidrug resistance protein 1

Disclosure; Page 54; 100pp; English.

polynucleotide

(EPID-) EPIDAUROS BIOTECHNOLOGIE AG.

Kerb R;

Heinrich G,

WPI; 2003-354397/33.

23-JUL-2002; 2002WO-EP008200.

WO2003013533-A2.

20-FEB-2003

Unidentified.

23-JUL-2001; 2001EP-00117608. 24-MAY-2002; 2002EP-00011710.

Sequence 19 BP; 6 A; 5 C; 4 G; 3 T; 0 U; 1 Other;

exemplification of the present invention

RESULT 1439

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ovarian cancer; pancreatic cancer; malignant glioma; uridine diphosphate glycosyltransferase1 member A1.
                                                          WO2003013536-A2
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                                    Homo sapiens.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     g
                                                                                                                                                                                                                                                                                                                                                                                                                                           The present invention describes a method for the use of irinotecan (I) or its derivative for the preparation of a pharmaceutical composition for treating colorectal, cervical, gastric, lung, ovarian or pancreatic cancer, or malignant glioma in a subject having a genome with a variant allele which comprises a multidrug resistance protein 1 (MRP1) polynucleotide (II). (I) has cytostatic activity. (I) or its derivative can be used for the preparation of a pharmaceutical composition for treating colorectal, cervical, gastric, lung, ovarian or pancreatic cancer, or malignant glioma in a subject, where the subject is a human (preferably African or Asian) or a mouse. The present sequence represents a sequence which is used in the exemplification of the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ö
                                                                                                     irinotecan, colorectal cancer, cervical cancer, gastric cancer, lung cancer, ovarian cancer, pancreatic cancer, malignant glioma, variant allele, multidrug resistance protein 1; MRP1; cytostatic, gene,
                                                                                                                                                                                                                                                                                                                                                              Use of irinotecan or its derivative for preparation of a pharmaceutical composition for treating cancer in a subject having a genome with a variant allele comprising a multidrug resistance protein 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ss; irinotecan; cancer; UGT1A1; cytostatic; topoisomerase I inhibitor; colorectal cancer; cervical cancer; lung cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human UGT1A1 variant allele sequence fragment SEQ ID NO:443.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 13.8; DB 1; Length 19; Pred. No. 9.8e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Indels
                                                                                MRP1 based cancer related nucleic acid SEQ ID NO:472.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 19 BP; 6 A; 5 C; 4 G; 3 T; 0 U; 1 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mismatches
                                                                                                                                                                                                                                                                                            (EPID-) EPIDAUROS BIOTECHNOLOGIE AG.
                                                                                                                                                                                                                                                                                                                                                                                                                         Disclosure; Page 54; 100pp; English.
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            ADB21314 standard; DNA; 19 BP.
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24-MAY-2002; 2002EP-00011710.
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                                                          (first entry)
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                                                                                                                                                                                                                                                                                                                    Kerb R;
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Best Local Similarity
                                                                                                                                                                                      WO2003013533-A2
                                                                                                                                                                                                                                                                                                                                                                                                   polynucleotide
                                                                                                                                                               Unidentified.
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                                                         20-NOV-2003
                                                                                                                                                                                                                                                                                                                    Heinrich G,
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 ADB21314/c
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Use of irinotecan to treat cancer patient by determining if patient has variant alleles of UGTIA1 gene, administering increased/decreased amounts of irinotecan based on increased/decreased levels of UGTIA1 gene product.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The invention relates to the novel use of irinotecan to treat a patient suffering from cancer. This involves determining if the patient has one or more variant alleles of the UGTA1 gene, and if the patient has one or more of such variant alleles, irinotecan is administered in an increased or decreased amount in comparison to the amount that is administered without regard to the patient's alleles in the UGTA1 gene. The invention has cytostatic activity. A composition of the invention acts as a topoisomerase I inhibitor. The method is useful for treating a patient,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         an animal e.g. mouse or a human, preferably African or Asian, suffering from cancer such as colorectal, cervical, gastric cancer, lung, ovarian pancreatic cancer or malignant glioma. The present sequence is udes in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ss; irinotecan; cancer; UGT1A1; cytostatic; topoisomerase I inhibitor;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 19;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 19 BP; 3 A; 4 C; 5 G; 6 T; 0 U; 1 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 13.8; DB 1;
Pred. No. 9.8e+02;
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                                                                                                                                                                                    (EPID-) EPIDAUROS BIOTECHNOLOGIE AG.
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23-JUL-2002; 2002WO-EP008217.
                                                                  23-JUL-2001; 2001EP-00117608.
24-MAY-2002; 2002EP-00011710.
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Best Local Similarity 78.9'
Matches 15, Conservative
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The invention relates to the novel use of irinotecan or its derivative
                                                                                the preparation of pharmaceutical compositions for treating
 multidrug resistance 1 polynucleotide.
                                  Disclosure; Page 82; 130pp; English.
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                                                                                                                                                                                                                                                The invention relates to the novel use of irinotecan to treat a patient suffering from cancer. This involves determining if the patient has one or more variant alleles of the UGTIAI gene, and if the patient has one or more of such variant alleles, irinotecan is administered in an increased or decreased amount in comparison to the amount that is administered without regard to the patient's alleles in the UGTIAI gene. The invention has cytostatic activity. A composition of the invention acts as a topoisomerase I inhibitor. The method is useful for treating a patient, an animal e.g. mouse or a human, preferably African or Asian, suffering from cancer such as colorectal, cervical, gastric cancer, lung, ovarian, phonoreatic or a human, preferably African or Asian, suffering from cancer such as colorectal, cervical, gastric cancer, lung, ovarian, the patient of the patient of the such as colorectal and the patient sequence is udes in
                                                                                                                                                Use of irinotecan to treat cancer patient by determining if patient has variant alleles of UCT1A1 gene, administering increased/decreased amounts of irinotecan based on increased/decreased levels of UGT1A1 gene product.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            irinotecan; colorectal cancer; cervical cancer; gastric cancer;
lung cancer; ovarian cancer; pancreatic cancer; malignant glioma;
multidrug resistance 1; MDR1; cytostatic; human; ds; Cyp3A5; MRP1; MDR1;
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                                                                                                                                                                                                                                                                                                                                                                                                                                      the exemplification of the invention.
                                                                                                                                                                                                                    Disclosure; Page 58; 107pp; English.
                                                  (EPID-) EPIDAUROS BIOTECHNOLOGIE AG.
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23-JUL-2001; 2001EP-00117608
24-MAY-2002; 2002EP-00011710
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nes 15; Conserv
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New use of irinotecan for preparation of pharmaceutical compositions for treating cancer in subject having genome with variant allele comprising

(EPID-) EPIDAUROS BIOTECHNOLOGIE AG.

WPI; 2003-268145/26.

Heinrich G,

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colorectal, cervical, gastric, lung, ovarian or pancreatic cancer, or malignant glioma in a subject having a genome with a variant allele which comprises a multidrug resistance I (MDRI) polynucleotide. A composition of the invention has cytostatic activity. The invention is useful for the preparation of pharmaceutical compositions for treating colorectal, cervical, gastric, lung, ovarian or pancreatic cancer, or malignant glioma in a subject (preferably human, more preferably African or Assian) or a mouse. The present sequence is used in the exemplification of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The invention relates to the novel use of irinotecan or its derivative for the preparation of pharmaceutical compositions for treating colorectal, cervical, gastric, lung, ovarian or pancreatic cancer, or malignant gliona in a subject having a genome with a variant allele which comprises a multidrug resistance 1 (MDR1) polynucleotide. A composition of the invention has cytostatic activity. The invention is useful for the preparation of pharmaceutical compositions for treating colorectal, cervical, gastric, lung, ovarian or pancreatic cancer, or malignant
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         lung cancer; ovarian cancer; pancreatic cancer; malignant glioma;
multidrug resistance 1; MDR1; cytostatic; human; ds; Cyp3A5; MRP1; MDR1;
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                                                                                                                                                                                                                                         Sequence 19 BP; 3 A; 4 C; 5 G; 6 T; 0 U; 1 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Disclosure; Page 82; 130pp; English.
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24-MAY-2002; 2002EP-00011710.
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ADB97386 standard; DNA; 19
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     04-DEC-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             20-FEB-2003
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SSSXS

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The invention relates ro a novel use of irinotecan or its derivative for the preparation of a pharmaceutical composition for treating colorectal, cervical, gastric, lung, ovarian or pancreatic cancer, or malignant glioma in a subject having a genome with a variant allele which comprises a multidrug resistance ! (MDR1) polynucleotide. A composition of the invention has cytostatic activity. The present sequence is used in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New use of irinotecan for preparation of pharmaceutical compositions for treating cancer in subject having genome with variant allele comprising multidrug resistance 1 polynucleotide.
                                                                                                                                                                irinotecan; colorectal cancer; cervical cancer; gastric cancer; lung cancer; ovarian cancer; pancreatic cancer; malignant glioma; multidrug resistance 1; MDR1; cytostatic; ds; human; UGT1A1; MRP1; TOP1.
                                                                                                                             Human MDR1 variant allele sequence fragment SEQ ID NO:472.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Hamster; high mobility group; HMG-I(Y); ss; primer; expression augmenting sequence element; EASE; RT-PCR; reverse transcriptase PCR; PCR.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Hamster high mobility group, HMG-I(Y), RT-PCR primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 19 BP; 6 A; 5 C; 4 G; 3 T; 0 U; 1 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0.8%; Score 13.8; DB 1; 78.9%; Pred. No. 9.8e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (EPID-) EPIDAUROS BIOTECHNOLOGIE AG.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Disclosure; Page 54; 104pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              exemplification of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               388 TCCTCGGATGAGGTGCAGT 406
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         19 TCCTCTGAGRATGTGCAGT 1
BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                       23-JUL-2001; 2001EP-00117608.
24-MAY-2002; 2002EP-00011710.
  ADB92577 standard; DNA; 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADD89803 standard; DNA; 19
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Kerb
                                                                                                                                                                                                                                                                                                WO2003013535-A2
                                                                                                                                                                                                                                                         Homo sapiens.
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                                                                                   04-DEC-2003
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                                         ADB92577;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The invention relates ro a novel use of irinotecan or its derivative for the preparation of a pharmaceutical composition for treating colorectal, cervical, gastric, lung, ovarian or pancreatic cancer, or malignant glioma in a subject having a genome with a variant allele which comprises a multidarug resistance I (WDR1) polymoclectide. A composition of the invention has cytostatic activity. The present sequence is used in the exemplification of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New use of irinotecan for preparation of pharmaceutical compositions for treating cancer in subject having genome with variant allele comprising multidrug resistance 1 polynucleotide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               irinotecan; colorectal cancer; cervical cancer; gastric cancer; lung cancer; ovarian cancer; pancreatic cancer; malignant glioma; multidrug resistance 1; MDR1; cytostatic; ds; human; UGT1A1; MRP1; TOP1.
subject (preferably human, more preferably African or Asian)
The present sequence is used in the exemplification of the
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                                                                                                                         Query Match 0.8%; Score 13.8; DB 1; Length 19; Best Local Similarity 78.9%; Pred. No. 9.8e+02; Matches 15; Conservative 1; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human MDR1 variant allele sequence fragment SEQ ID NO:471.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 19 BP; 3 A; 4 C; 5 G; 6 T; 0 U; 1 Other;
                                                                                   Sequence 19 BP; 6 A; 5 C; 4 G; 3 T; 0 U; 1 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (EPID-) EPIDAUROS BIOTECHNOLOGIE AG.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Disclosure; Page 54; 104pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           388 TCCTCGGATGAGGTGCAGT 406
                                                                                                                                                                                                               388 TCCTCGGATGAGGTGCAGT 406
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                                                                                                                                                                                                                                                      19 rccrcrcagrarcrcaci
                                                                                                                                                                                                                                                                                                                                                              BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                23-JUL-2002; 2002WO-EP008220
                                                                                                                                                                                                                                                                                                                                                              ADB92576 standard; DNA; 19
                                                                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Kerb R;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2003-342400/32.
    glioma in a subject
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                                                                                                                                                                                                                                                                                                                                                                                                                                             04-DEC-2003
                        or a mouse.
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ADB92576;

RESULT 1444 ADB92576

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Gaps

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3; Indels

US2003008345-A1

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RESULT 1445 ADB92577/c

09-JAN-2003

Length 19;

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                                                                                                                                                                                                                                                                                                                                                                     The invention relates to an isolated polynucleotide comprising a nucleic acid molecule comprising nucleotides 11538-11692, nucleotides 11538-1170, nucleotides 11673-12165, nucleotides 11813-12165 or nucleotides 11899-12165 of ADD89798, the hamster high mobility group, HWG-I(Y) gene, fragments of the DNA having expression augmenting activity (an expression augmenting sequence element, EASE) or their combinations or complementary DNA. Also included are a mammalian host cell which comprises the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          short interfering nucleic acid; siNA; downregulation; inhibition; SCD; stearcyl-CoA desaturase; RNA interference; anorectic; antidiabetic; antiarteriosclerotic; cytostatic; virucide; obesity; diabetes; atherosclerosis; cancer; viral infection; drug screening;
                                                                                                                                                                                                                                                                                   New isolated polynucleotide used for producing recombinant protein by
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
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88.2%; Pred. No. 9.8e+02;
ative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Stearoyl-CoA desaturase siNA oligonucleotide SEQ ID NO:462.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   genetic engineering; pharmacogenomic; gene mapping; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 19 BP; 2 A; 9 C; 3 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                       Example 16; Page 13; 27pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           390 CTCGGATGAGGTGCAGT 406
                                                                                                                                                                                                                                                                                                    culturing mammalian host cell.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADE27518 standard; RNA; 19 BP
                                            11-JAN-1996, 96US-00586509.
13-JAN-1997; 97US-00785150.
0S-NOV-1999; 99US-0043377.
02-MAR-2000; 2000US-0186537P.
12-SEP-2000; 2000US-00660299.
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              09-OCT-2001; 2001US-00973928
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Best Local Similarity 88.2
Matches 15; Conservative
                                                                                                                                                                                                            Thomas JN;
                                                                                                                                                                                                                                              WPI; 2003-863362/80.
                                                                                                                                                        (MORR/) MORRIS A E. (THOM/) THOMAS J N.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WO2003070885-A2
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                                                                                                                                                                                                            Morris AE,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          short interfering nucleic acid; siNA; downregulation; inhibition; SCD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   studying gene function and gene mapping (e.g. of single-nucleotide polymorphisms). The present sequence represents an SCD siNA, which is used in the exemplification of the present invention.
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                                                                                                                                                                                                                                                                  diagnosis of obesity or diabetes, downregulates expression of the stearcyl-CoA desaturase gene.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 19 BP; 5 A; 3 C; 8 G; 0 T; 3 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                Example 3; SEQ ID NO 462; 139pp; English.
                                                                                                                                                                                                          Thompson J;
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          11-MAR-2002; 2002US-0363124P.
06-JUN-2002; 2002US-0386782P.
29-AUG-2002; 2002US-0406784P.
05-SEP-2002; 2002US-0409378P.
09-SEP-2002; 2002US-040923P.
20-SEP-2002; 2002US-0412304P.
15-JAN-2003; 2003US-0440129P.
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                                                                                                                                                                                                        Mcswiggen J, Beigelman L,
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ID ADE27228 standard; RNA; 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2 AGGUGGAGACACUGCGG
                                                                                                                                                                    (RIBO-) RIBOZYME PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first entry)
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nucleic acid (siNA) that downregalates expression of the vascular and nucleic acid (siNA) that downregalates expression of the vascular endothelial growth factor receptor (VEGFR) gene. Also described: (1) a siNA that downregulates the VEGF gene; (2) kits for in vitro or in vivo delivery of siNA; (3) conjugates and/or complexes of siNA; (4) vectors that express siNA; and (5) single-stranded siNA with similar properties. The siNAs have antiangiogenic, cytostatic, nephrotropic and gynaecological, antiarthritic, antipsoriatic, nephrotropic and gynaecological activities. The siNA are useful for modulating useful for treating wide range of angiogeness rassociated conditions, particularly cancers, diabetic retinopathy, macular degeneration, and polycystic kidney disease. The siNA may also be useful for diagnosis, and polycystic kidney disease. The siNA may also be useful for diagnosis, druging studying gene function, and validation, genetic engineering, studying gene function, and also for gene mapping (e.g. of exemplification of the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         double-stranded short interfering nucleic acid; short interfering nucleic acid; siNA; downregulation; vascular endothelial growth factor receptor; VEGFR; antiangiogenic; cytostatic; antidiabetic; ophthalmological; antiarthritic; antipsoriatic; nephrotropic; gynaecological; angiogenesis-associated condition; cancer; diabetic retinopathy; macular degeneration; neovascular glaucoma;
                                                                                                                                                                                                                                                                                                                                                                                                                      New double-stranded interfering nucleic acid, useful e.g. for treatment and diagnosis of cancer, downregulates the vascular endothelial growth factor receptor gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The present invention describes a double-stranded short interfering
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human VEGFR3 short interfering nucleic acid (sinA) SEQ ID NO:1792.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 19 BP; 5 A; 5 C; 5 G; 0 T; 4 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 3; SEQ ID NO 1545; 207pp; English.
                                                                                                                                                                                                                                                                                                                                                  Beigelman L, Pavco P;
                                                     11-MAR-2002; 2002US-0363124P.
29-MAY-2002; 2002WO-US017674.
03-JUL-2002; 2002US-0396782P.
03-JUL-2002; 2002US-0399348P.
29-JUL-2002; 2002US-0399348P.
29-AUG-2002; 2002US-0406784P.
05-SEP-2002; 2002US-0406784P.
05-SEP-2002; 2002US-040639978P.
04-NOV-2002; 2002US-0409293P.
27-NOV-2002; 2002US-04092949.
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20-FEB-2003; 2003WO-US005022
                                      2002US-0358580P.
                                                                                                                                                                                                                                                                   15-JAN-2003; 2003US-0440129P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          76.5%;
                                                                                                                                                                                                                                                                                                         (RIBO-) RIBOZYME PHARM INC.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 12-FEB-2004 (first entry)
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Matches 13; Conservative
                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2003-679876/64.
                                                                                                                                                                                                                                                                                                                                                  Mcswiggen J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADF37503;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     double-stranded short interfering nucleic acid; short interfering nucleic acid; siNA; downregulation; vascular endothelial growth factor receptor; VEGFR; antiangiogenic; cytostatic; antidiabetic; ophthalmological; antiarthritic; antipsoriatic; nephrotropic; gynaecological; angiogenesis-associated condition; cancer; diabetic retinopsthy; macular degeneration; neovascular glaucoma; arthritis; psoriasis; endometriosis; angiofibroma;
                                                                                                                                                                                                                                                                                                                                                                                                       The present invention describes a short interfering nucleic acid (siNA) that downregulates expression of the SCD (stearoyl-CoA desaturase) gene by RNA interference. Also described: (1) modulating expression of SCD genes in cells, tissue explants or organisms by introduction of StD sins fits for in vitro or in vivo delivery of siNA, (3) conjugates and/or complexes of siNA, and (4) vectors that express siNA. SCD inhibiting siNAs have ancrectic, antidiabetto, antiatreriosclerotic, crytostatic and virucide activities. The siNAs can be used to modulate expression of SCD genes, in cells, tissue explants or organisms, e.g. for treating obesity, diabetes (types I and II); atherosclerosis; cancer and viral infections. They can also be used for drug screening; diagnosis; target studying gene function and validation; genetic engineering; pharmacogenomics; studying gene function and gene mapping (e.g. of single-nucleotide polymorphisms). The present sequence represents an SCD siNA, which is
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human VEGFR3 short interfering nucleic acid (siNA) SEQ ID NO:1545.
                                                                                                                                                                                                                                                                                    New short interfering nucleic acid, useful e.g. for treatment and diagnosis of obesity or diabetes, downregulates expression of the stearoyl-CoA desaturase gene.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    used in the exemplification of the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 19 BP; 3 A; 8 C; 3 G; 0 T; 5 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mismatches
                                                                                                                                                                                                                                                                                                                                                                   Example 3; SEQ ID NO 172; 139pp; English.
                                                                                                                                                                                                     Mcswiggen J, Beigelman L, Thompson J;
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                                      29-AUG-2002; 2002US-0406784P.
05-SEP-2002; 2002US-0408378P.
09-SEP-2002; 2002US-0409237P.
20-SEP-2002; 2002US-0412304P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADF37256 standard; RNA; 19 BP.
2002US-0363124P.
2002US-0386782P.
                                                                                                                         15-JAN-2003; 2003US-0440129P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          18 AGGTGGAGACACTGCGG 2
                                                                                                                                                               (RIBO-) RIBOZYME PHARM INC
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Best Local Similarity 88.2
Matches 15; Conservative
                                                                                                                                                                                                                                              WPI; 2003-721687/68.
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11-MAR-2002;
06-JUN-2002;
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Gaps

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10017621-381.rng

RNA interference; short interfering nucleic acid; siNA; short interfering RNA; siRNA; double-stranded RNA; micro-RNA; miRNA; short hairpin RNA; shRNA; expression modulation; gene therapy;

Human IGF-1R siNA lower strand, SEQ ID NO:370.

12-FEB-2004 (first entry)

drug screening; diagnosis; therapeutic target identification; pharmacogenomics; gene function analysis; gene mapping; cancer; proliferative disease; restenosis; polycystic kidney disease; inflammatory disease; allergic disease; autoimmune disease; transplant rejection; cytostatic; vasotropic; nephrotropic; antiinflammatory; antiallergic; immunosuppressive; human; insulin-like growth factor I receptor; IGF-IR; ss.

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nucleic acid (siNN) that downregulates expression of the vascular candchelial growth factor receptor (VEGFR) gene. Also described: (1) a siNA that downregulates the VEGF gene; (2) kits for in vitro or in vivo delivery of siNA; (3) conjugates and/or complexes of siNA; (4) vectors that express siNA; and (5) single-stranded siNA with similar properties. The siNAs have antiangiogenic, cytostatic, antidiabetic, ophthalmological, antiathritic, antigoriatic, nephrotropic and gynaecological, antiathritic, antigoriatic, nephrotropic and gynaecological activities. The siNA are useful for modulating (downregulating) the expression of VEGFR genes. The siNA are potentially useful for treating wide range of angiogenesis-associated conditions, particularly cancers, diabetic retinopathy, macular degeneration, particularly cancers, diabetic retinopathy, macular degeneration, and polycystic kidney disease. The siNA may also be useful for diagnosis, and polycystic kidney disease. The sinA may also be useful for diagnosis, chung screening, target identification and validation, genetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New double-stranded interfering nucleic acid, useful e.g. for treatment and diagnosis of cancer, downregulates the vascular endothelial growth
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in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The present invention describes a double-stranded short interfering
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                and also for gene mapping e present sequence is used
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                engineering, studying gene function, and also for gene map
single-nucleotide polymorphisms). The present sequence is
endometriosis; angiofibroma;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 19 BP; 4 A; 5 C; 5 G; 0 T; 5 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mcswiggen J, Beigelman L, Pavco P;
                                                                                                                                                                                                                                                                                         2002US-0393796F
2002US-0399348P
2002US-0406784P
2002US-0408378P
2002US-0409293P
2002US-04087949.
arthritis; psoriasis; endomet
polycystic kidney disease; ss
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2002WO-US017674.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                               (RIBO-) RIBOZYME PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               factor receptor gene.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      exemplification of
                                                                                                           WO2003070910-A2.
                                                                          Homo sapiens
                                                                                                                                                                                   20-FEB-2003;
                                                                                                                                                                                                                                                                                           03-JUL-2002;
                                                                                                                                                                                                                                                                                                                                                    05-SEP-2002;
                                                                                                                                                                                                                                                                                                                                                                     09-SEP-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                           15-JAN-2003;
                                                                                                                                                                                                                                                                           06-JUN-2002;
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                                                                                                                                                                                                                        20-FEB-2002;
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                                                                                                                                                                                                                                         11-MAR-2002
29-MAY-2002
                                                                                                                                               28-AUG-2003
                                                     Synthetic
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Mcswiggen J, Beigelman L, Chowrira B;

WPI; 2003-721691/68

(RIBO-) RIBOZYME PHARM INC.

; 2002US-0406784P. ; 2002US-0408378P. ; 2002US-0409293P. ; 2003US-0440129P.

29-AUG-2002; 05-SEP-2002; 09-SEP-2002;

L5-JAN-2003;

2002US-0363124P. 2002US-0386782P.

20-FEB-2003; 2003WO-US005044.

20-FEB-2002; 11-MAR-2002; 06-JUN-2002;

WO2003070911-A2.

28-AUG-2003.

Homo sapiens

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New short interfering nucleic acid, useful e.g. for treatment and diagnosis of cancer , downregulates expression of the insulin-like growth
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The invention relates to short interfering nucleic acids (siNA) which downregulate expression of the human insulin-like growth factor 1 receptor (IGF-IR) gene by RNA interference. The siNAs may or may not comprise ribonucleotides and may be double or single stranded. They further comprise sense and antisense regions, or alternatively are assembled from a sense oligonucleotide and an antisense oligonucleotide Specifically, the siNAs include short interfering RNA (siRNA), doublestranded RNA, micro-RNA (miRNA) and short hairpin RNA (shRNA). The siNAs can be unmodified or chemically modified, can contain
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               mapping (e.g., of single nucleotide polymorphisms). The present sequence represents the lower strand of a human IGF-1R-targeted double-stranded
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              genetic engineering, pharmacogenomics, studying gene function, and gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 19 BP; 4 A; 8 C; 4 G; 0 T; 3 U; 0 Other;
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nes 13; Conservative
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Best Local Si
Matches 13;
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Gaps

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0.8%; Score 13.8; DB 1; Length 19; 38.2%; Pred. No. 9.8e+02; ive 0; Mismatches 2; Indels

88.2%;

Conservative

Local Similarity ses 15; Conserv

Best Loca Matches

Query Match

517 GAGAAGCTGACCCTCAA 533

GAGAAGCTGGTCCTCAA

ADF31705 standard; RNA; 19 BP

RESULT 1451

ADF31705

ADF31705;

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further comprises eachse and antisense regions, or alternatively are assembled from a sense oligomoclectide and an antisense oligomoclectide and an antisense oligomoclectide. Specifically, the sinAs include short interfering RNA (siRNA), doublestranded RNA, micro-RNA (miRNA) and short hairpin RNA (siRNA). The siNAs can be unmodified or chemically modified, can contain decoxyribomoclectides, and can be chemically synthesised, expressed from a vector or enzymatically synthesised. The invention also relates to kits for the in vitro or in vivo delivery of siNA; conjugates and/or complexes of siNA; and vectors that express siNA. The siNAs are used to modulate expression of the IGF-IR gene in cells, tissue explants or organisms (e.g., by ex vivo gene therapy), or in grafts and transplants for the transmittery of a variety of conditions. They may be used for treating cancer and other proliferative diseases (e.g., restenosis and polycystic kidney disease), inflammatory and/or allergic diseases, autoimmune
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New short interfering nucleic acid, useful e.g. for treatment and diagnosis of cancer , downregulates expression of the insulin-like growth
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human IGF-1R transcript target sequence/siNA upper strand, SEQ ID NO:93.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RNA interference; short interfering nucleic acid; siNA; short interfering RNA; siRNA; double-stranded RNA; micro-RNA; miRNA; short interfering RNA; shRNA; expression modulation; gene therapy; and arrows a screening; diagnosis; therapeutic target identification; pharmacogenomics; gene function analysis; gene mapping; cancer; proliferative disease; restenosis; polycystic kidney disease; inflammatory disease; allergic disease; autoimmune disease; transplant rejection; cytostatic; vasotropic; nephrotropic; antiinflammatory; antiallergic; immunosuppressive; human;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     insulin-like growth factor 1 receptor; IGF-1R; target sequence; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example 3; SEQ ID NO 93; 147pp; English.
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1299
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2002US-0363124P.
2002US-0386782P.
2002US-0406784P.
2002US-0409378P.
2002US-0409233P.
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CAGGCATCCTGTCCAAC
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                                                                                                                                                                                                                                                             ADF31428 standard; RNA; 19
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Homo sapiens.
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29-AUG-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                05-SEP-2002;
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11-MAR-2002;
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1283
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                                                                                                                                                                                       RESULT 1452
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 for prognosticating, diagnosing and treating cardiovascular diseases, such as atherosclerosis, hypertension, restenosis, arterial inflammation, myocardial infarction and stroke. The present sequence is a PCR primer, used in the examples from the invention.
                                 genetic engineering, pharmacogenomics, studying gene function, and gene mapping (e.g., of single nucleotide polymorphisms). The present sequence represents the upper strand of a human IGF-IR-targeted double-stranded siNA, which is identical to the IGF-IR transcript target sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Cardiant; antiarteriosclerotic; vasotropic; cerebroprotective; hypotensive; gene therapy; human; apolipoprotein C-III; PCR; primer; ss.
diseases and transplant rejection. The siNAs are also useful for drug screening, diagnosis, therapeutic target identification and validation,
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                                                                                                                                                                                          Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New isolated polynucleotide encoded by a phenotype-associated gene, useful for prognosticating statin therapy response, and diagnosing treating cardiovascular diseases, such as hypertension, myocardial
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        .;
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                                                                                                                                                        Length 19;
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88.2%; Pred. No. 9.8e+02;
ive 0; Mismatches 2; Indels
                                                                                                                                                                                          Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 19 BP; 2 A; 10 C; 2 G; 5 T; 0 U; 0 Other;
                                                                                                                          BP; 3 A; 4 C; 8 G; 0 T; 4 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Apolipoprotein C-III, BaySNP 1837, PCR primer #2.
                                                                                                                                                                                          7
                                                                                                                                                        Score 13.8; DB 1;
Pred. No. 9.8e+02;
                                                                                                                                                                                        0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 1; Page 67; 182pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Kallabis
                                                                                                                                                                                                                            1283 CAGGCATCCTGTCCAAC 1299
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            306 CCCACTCAGCTCTGCAC 322
                                                                                                                                                                                                                                                                                                                                                 BP
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                                                                                                                                                            0.88;
                                                                                                                                                                          88.2%;
                                                                                                                                                                                                                                                              18 CAGGCATCCTGCCCATC
                                                                                                                                                                                                                                                                                                                                                 13
                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
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Best Local Similarity 88.2
Matches 15; Conservative
                                                                                                                                                                                              Conservative
                                                                                                                                                                                                                                                                                                                                                ADF13396 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ຜ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 infarction and stroke.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2003-712738/67
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Schwers
                                                                                                                                                                          Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (FARB ) BAYER AG.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WO2003072813-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Homo sapiens
                                                                                                                          Sequence 19
                                                                                                                                                                                                                                                                                                                                                                                                                 12-FEB-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          04-SEP-2003.
                                                                                                                                                                                              15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Stropp U,
                                                                                                                                                                                                                                                                                                                                                                                 ADF13396;
                                                                                                                                                            Query Match
                                                                                                                                                                                                                                                                                                              RESULT 1453
                                                                                                                                                                                              Matches
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ADF84791;

1 CCCACTCAGCCCTGCTC 17

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The invention relates to a novel double-stranded short interfering nucleic acid (siNA) that downregulates expression of the breakpoint cluster region-vabl Abelson murine leukaemia viral oncogene homologue 1 (BCR-ABL) gene. The siRNA of the invention demonstrates cytostatic activity and may be useful for modulating expression of the BCR-ABL gene, as well as for treating leukaemia or lymphoma and in diagnosis, drug screening, target identification and validation, genetic engineering, gene function studies and gene mapping. The current sequence is that of the human ABL1-targeted siRNA of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New double-stranded interfering nucleic acid, useful e.g. for treatment and diagnosis of leukemia and lymphoma, downregulates the breakpoint
                                                                                                                                               short interfering nucleic acid; siNA; breakpoint cluster region; v-abl Abelson murine leukaemia viral oncogene homologue 1; BCR-ABL; cytostatic; leukaemia; lymphoma; human; ss; siRNA; ABL1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0.8%; Score 13.8; DB 1; Length 19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 19 BP; 1 A; 8 C; 4 G; 0 T; 6 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Pred. No. 9.8e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example 7; SEQ ID NO 921; 197pp; English.
                                                                                                                  Human ABL1-targeted siRNA - SEQ ID 921
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Chowrira B;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         and diagnosis of leukemia and lymphoma
cluster region-Abelson (BCR-ABL) gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ;
                                                                                                                                                                                                                                                                                                                                               20-FEB-2002; 2002US-0358580P.

11-MAR-2002; 2002US-0363124P.

06-JUN-2002; 2002US-0366782P.

15-AUG-2002; 2002US-0404039P.

29-AUG-2002; 2002US-0406784P.

05-SEP-2002; 2002US-0408378P.

09-SEP-2002; 2002US-040923P.

14-JAN-2003; 2003US-0439922P.

15-JAN-2003; 2003US-0439922P.
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                                                                                                                                                                                                                                                                                                                   20-FEB-2003; 2003WO-US005234
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        88.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Beigelman L,
               ADF84627 standard; RNA; 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (RIBO-) RIBOZYME PHARM INC
                                                                                 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2003-679889/64.
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                                                                                                                                                                                                                                                   WO2003070972-A2
                                                                                                                                                                                                                    Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mcswiggen J,
                                                                                 26-FEB-2004
                                                                                                                                                                                                                                                                                    28-AUG-2003
                                                  ADF84627;
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The invention relates to a novel double-stranded short interfering nucleic acid (siNA) that downregulates expression of the breakpoint cluster region-v-abl Abelson murine leukaemia viral oncogene homologue 1 (BCR-ABL) gene. The siRNA of the invention demonstrates cytostatic activity and may be useful for modulating expression of the BCR-ABL gene, as well as for treating leukaemia or lymphoma and in diagnosis, drug screening, target identification and validation, genetic engineering, gene function studies and gene mapping. The current sequence is that of the human ABL1-targeted siRNA of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New double-stranded interfering nucleic acid, useful e.g. for treatment and diagnosis of leukemia and lymphoma, downregulates the breakpoint cluster region-Abelson (BCR-ABL) gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
                                                                                      short interfering nucleic acid; siNA; breakpoint cluster region; v-abl Abelson murine leukaemia viral oncogene homologue 1; BCR-ABL; cytostatic; leukaemia; lymphoma; human; ss; siRNA; ABL1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       short interfering nucleic acid; siNA; breakpoint cluster region; v-abl Abelson murine leukaemia viral oncogene homologue 1; BCR-ABL;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 19 BP; 5 A; 5 C; 8 G; 0 T; 1 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 7; SEQ ID NO 1085; 197pp; English.
                                                            fuman ABL1-targeted siRNA - SEQ ID 1085.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Chowrira B;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human ABL1-targeted siRNA - SEQ ID 602.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GGGCAGAGUGACCAGCC 18
                                                                                                                                                                                                                                                                                                              2002US-0386782P.
2002US-0404039P.
2002US-0406784P.
                                                                                                                                                                                                                                                                                                                                                          05-SEP-2002; 2002US-0408378P.
09-SEP-2002; 2002US-0409293P.
14-JAN-2003; 2003US-0439922P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADF84308 standard; RNA; 19
                                                                                                                                                                                                                                                                                                                                                                                                                                        (RIBO-) RIBOZYME PHARM INC
                             (first entry)
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                                                                                                                                                                                   WO2003070972-A2
                                                                                                                                                                                                                                                                                             11-MAR-2002;
06-JUN-2002;
                                                                                                                                                       Homo sapiens.
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                             26-FEB-2004
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Best Local S
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Gaps

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ADF84791 standard; RNA; 19 BP

RESULT 1455

ADF84791 ID ADF

GCAACCGGCAGGAGGTG

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SNP; single nucleotide polymorphism; cardiovascular associated gene; allelic variation; atherosclerosis; ischemia; reperfusion; hypertension; restenosis; arterial inflammation; myocardial infarction; stroke; primer;
                                                                                                                                                                                                                                                                                                                                                                                                                               The invention relates to a novel double-stranded short interfering nucleic acid (siNA) that downregulates expression of the breakpoint cluster region-v-abl Abelson murine leukaemia viral omoogene homologue 1 (BCR-ABL) gene. The siRNA of the invention demonstrates cytostatic activity and may be useful for modulating expression of the BCR-ABL gene, as well as for treating leukaemia or lymphoma and in diagnosis, drug screening, target identification and validation, genetic engineering, gene function studies and gene mapping. The current sequence is that of the human ABL1-targeted siRNA of the invention.
                                                                                                                                                                                                                                                                                                                         New double-stranded interfering nucleic acid, useful e.g. for treatment and diagnosis of leukemia and lymphoma, downregulates the breakpoint cluster region-Abelson (BCR-ABL) gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0.8%; Score 13.8; DB 1; Length 19; 38.2%; Pred. No. 9.8e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 19 BP; 1 A; 8 C; 5 G; 0 T; 5 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0; Mismatches
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                                                                                                                                                                                                                                                       Chowrira B;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Kallabis H;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                361 GGGGAGAGTGACCAGGC 377
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              BP.
                                 2002US-0358580P.
2002US-0363124P.
2002US-0386782P.
                                                                                    15-AUG-2002; 2002US-0404039P.
29-AUG-2002; 2002US-0406784P.
05-SEP-2002; 2002US-0406787PP.
09-SEP-2002; 2002US-0408237P.
14-UAN-2003; 2003US-0439922P.
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20-FEB-2003; 2003WO-US005234
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                                                                                                                                                                                                                                                       Beigelman L,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  18 GGCCAGAGTGACCAGCC
                                                                                                                                                                                                                     (RIBO-) RIBOZYME PHARM INC
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                                                                                                                                                                                                                                                                                           WPI; 2003-679889/64.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (FARB ) BAYER AG
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                                                                                                                                                                                                                                                         Mcswiggen J,
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                                                      11-MAR-2002;
06-JUN-2002;
                                     20-FEB-2002;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The invention relates to a novel double-stranded short interfering nucleic acid (siNA) that downregulates expression of the breakpoint oluster region-v-abl Abelson murine leukaemia viral oncogene homologue 1 (BCR-ABL) gene. The siNNA of the invention demonstrates cycostatic activity and may be useful for modulating expression of the BCR-ABL gene, as well as for treating leukaemia or lymphoma and in diagnosis, drug screening, target identification and validation, genetic engineering, gene function studies and gene mapping. The current sequence is that of the human ABL1-targeted siRNA of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New double-stranded interfering nucleic acid, useful e.g. for treatment and diagnosis of leukemia and lymphoma, downregulates the breakpoint cluster region-Abelson (BCR-ABL) gene.
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     cytostatic; leukaemia; lymphoma; human; ss; siRNA; ABL1.
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09-SEP-2002; 2002US-U4ve-2...
09-SEP-2002; 2002US-0409293P.
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2002US-0406784P.
2002US-0408378P.
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2002US-0386782P.
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                                                                                                                                                                                                                                                                                                                                                                      (RIBO-) RIBOZYME PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                        Mcswiggen J, Beigelman L,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 2003-679889/64.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WO2003070972-A2.
                                                                             WO2003070972-A2
                                                                                                                                                                                                                                        15-AUG-2002;
29-AUG-2002;
                                                                                                                                                                                                                                                                                                                                  15-JAN-2003;
                                                                                                                                                                                                      11-MAR-2002;
06-JUN-2002;
                                                                                                                                                                                       20-FEB-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Homo sapiens
                                           Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   26-FEB-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     28-AUG-2003
                                                                                                                 28-AUG-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADF84472;
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Matches

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Gaps

0

Indels

US2003232437-A1 Zhang H, RESULT 1459 ADH54705, 8 a

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The invention relates an isolated polymucleotide (I) encoded by a cardiovascular associated (CA) gene, having allelic variation contained in a functional surrounding like full length cDNA for CA gene polypeptide, and with or without the CA gene promoter sequence. (I) is a polymucleotide comprising single nucleotide polymorphisms predicting ardiovascular disease. The polymucleotide are useful for assessing predisposition or susceptibility to a cardiovascular disease, e.g. attherosolerosis, ischemia/reperfusion, hypertension, restenosis, arterial inflammation, myocardial infarction, and stroke. These may also be used to predict personal medication schemes omitting adverse drug reactions, or as probes for detecting genetic polymorphisms and as templates for the recombinant production of normal or variant peptides/polypeptides encoded by the genes. This sequence corresponds to a PCR primer to amplify one of
                                                                                                    New isolated polynucleotides comprising single nucleotide polymorphisms of the cardiovascular gene, useful for assessing predisposition or susceptibility to a cardiovascular disease, e.g. atherosclerosis,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 19 BP; 2 A; 10 C; 2 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                               Disclosure; Page 77; 187pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          the genes of the invention.
                                                                                                                                                                                                                                                                                     restenosis or stroke.
WPI; 2003-577532/54.
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Gaps
                                 0;
  Score 13.8; DB 1; Length 19;
Pred. No. 9.8e+02;
0; Mismatches 2; Indels
                               .
0
                                                           322
                                                                                    1 cccacrcacccractc 17
  0.8%;
                                                          306 CCCACTCAGCTCTGCAC
Query Match 0.8
Best Local Similarity 88.2
Matches 15; Conservative
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0

ADH54705 standard; DNA; 19 Human VEGF-C PCR primer #2. 25-MAR-2004 (first entry) Homo sapiens ADH54705;

human; ss; PCR; VEGF-C; cardiovascular disorder; atherosclerosis; diabetic retinopathy; autoimmune disorder; inflammatory disorder; vascular endothelial growth factor; primer.

18-DEC-2003

17-JUN-2002; 2002US-00173718

17-JUN-2002; 2002US-00173718.

(ISIS-) ISIS PHARM INC

Dobie KW;

WPI; 2004-061284/06

The invention describes an isolated polypeptide (I) comprising a cids as given in the specification or a region consisting of 5 or more contiguous amino acids, where the region includes amino acid 6 634 of (S1). Also described are: an isolated polymuclectide (II) comprising a fully defined sequence (S2) of 2405 base pair as given in the specification, or its complement, a polymuclectide (II) comprising a specification, or its complement, a polymuclectide, or a region of 15 or more contiguous nucleotides, the region comprising nucleotide 1904 of (S2); vector, preferably an expression vector (III) comprising (II); a host cell (IV) comprising (II); a chost cell (IV) comprising (II); a chost cell (IV) comprising (II); detecting obesity or obesity.

The lated disorders associated with differential expression of a LIPIN3 polymucleotide comprising a detecting a level of expression of (V), or (VI), or a region of (V) or (VI), where the region is 10 or more concluded and of (V) or (VI), where the region is 10 or more concluded (II); and to (II); a recombinant cell sample; antibodies that specifically bind to (I); a recombinant cell sample; antibodies that specifically bind to (I); a recombinant cell somprising a recombinantly modified (II); and array comprising two or more (II); and identifying an alteration in

New compounds, particularly antisense oligonucleotides targeted to a nucleic acid encoding vascular endothelial growth factor-C (VEGF-C), useful for treating atherosclerosis, diabetic retinopathy, or inflammatory disorders.

Example 13; SEQ ID NO 6; 83pp; English

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The invention relates to a compound targeted to and which specifically hibridises with a nucleic acid molecule encoding VBGF-C, and inhibits the expression of VBGF-C. The compound, composition and methods are useful for treating a disease or condition associated with VBGF-C, such as a cardiovascular disorder e.g. atherosclerosis or diabetic retinopathy or an autoimmune or inflammatory disorder. They are also useful in research and diagnostics for modulating the expression of VBGF-C. The present sequence represents a human VBGF-C PCR primer.
                                                                                                                                                                                  Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                 LIPIN3; obesity; obesity-related disorder; differential expression; polynucleotide polymorphism; adipocyte; human; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Novel isolated LIPIN3 polypeptide, useful for diagnosing diabetes
                                                                                                                                                                                 ;
                                                                                                                                                    Query Match 0.8%; Score 13.8; DB 1; Length 19; Best Local Similarity 88.2%; Pred. No. 9.8e+02; Matches 15; Conservative 0; Mismatches 2; Indels
                                                                                                                          Sequence 19 BP; 3 A; 6 C; 5 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                        Human LIPIN3 exon14 PCR primer segid 42.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example 5; SEQ ID NO 41; 58pp; English.
                                                                                                                                                                              0;
                                                                                                                                                                                                          1563 GATGCCTGACTCAGGCA 1579
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          26-JUL-2002; 2002US-00206618.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    26-JUL-2002; 2002US-00206618.
                                                                                                                                                                                                                                    17 GATGCCTGGCTCAGGAA
                                                                                                                                                                                                                                                                                                      ADO17050 standard; DNA; 19
                                                                                                                                                                                                                                                                                                                                                            (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2004-122019/12.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (WARD/) WARDEN C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     US2004018497-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                             01-JUL-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               29-JAN-2004.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Warden CH;
                                                                                                                                                                                                                                                                                                                                  ADO17050;
                                                                                                                                                                                                                                                                               RESULT 1460
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is useful for detecting a polynucleotide polymorphism associated with obesity. (II) is useful for diagnosing obesity an obesity-related with disorder which involves detecting (I). In (M2), the cell is an adipocyte. The test agent is chosen from antibody, protein, mucleic acid, and small organic molecule. This sequence represents a primer used to identify single nucleotide polymorphisms in the human Lipin3 gene that may be
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Selecting siRNA by selecting an siRNA molecule of 19-25 nucleoside bases by selecting a target gene and measuring the functionality of the nucleotide sequences that are complementary to a stretch of nucleotides
                                                                                                                                             Gaps
                                                                                                                                                                                                                                                                                                                             Bcl-2; optimised; short interfering RNA; luciferase.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The invention relates to a novel method for selecting siRNA (short
                                                                                                                                             .;
0
                                                                                                                       0.8%; Score 13.8; DB 1; Length 19; 88.2%; Pred. No. 9.8e+02;
                                                                                                                                             Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Stephen
                                                                                                                                                                                                                                                                                                           Anti-Firefly luciferase siRNA Luc 78 SEQ ID NO:288
                                                                                                   Sequence 19 BP; 5 A; 9 C; 2 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          William M,
                                                                                                                                             Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 1; SEQ ID NO 288; 199pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Devin L,
                                                                                                                                             .;0
                                                                                                                                                                  1399
                                                                                                                                                                                        19
                                                                                                                                                                                                                                             BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                      14-NOV-2002; 2002US-0426137P.
                                                                                                                                                                                                                                                                                                                                                                                                                                 14-NOV-2003; 2003WO-US036787
                                                                                                                                                                                                                                                                                                                                                                                                                                                                10-SEP-2003; 2003US-0502050P
                                                                                                                                                                  1383 CGACCTCCTCACCAAGC
                                                                                                                                                                                        3 CGACCACTTCACCAAGC
                                                                                                                                                                                                                                                                                                                                gene silencing;
                                                                                                                                                                                                                                            ADQ60589 standard; RNA; 19
                                                                                                                                                                                                                                                                                                                                             RNA interference; firefly;
                                                                                                                                              Conservative
                                                                               associated with obesity.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ď
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DHAR-) DHARMACON INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Angela
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2004-420527/39.
                                                                                                                                                                                                                                                                                      (first
                                                                                                                                Local Similarity
nes 15; Conserv
                                                                                                                                                                                                                                                                                                                                                                                       WO2004045543-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      the target
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Anastasia K,
                                                                                                                                                                                                                                                                                      09-SEP-2004
                                                                                                                                                                                                                                                                                                                                                                                                            03-JUN-2004
                                                                                                                                                                                                                                                                                                                               ss; siRNA;
                                                                                                                                                                                                                                                                   ADQ60589;
                                                                                                                         Query Match
                                                                                                                                                                                                                        RESULT 1461
                                                                                                                                              Matches
                                                                                                                                                                                                                                   ADQ60589,
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       pairs. The kit comprises at least two siRNA, comprising a first optimised siRNA and a second optimised siRNA. The method is useful in selecting siRNA for generating a gene silencing reagent. The present sequence is used in the exemplification of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Selecting siRNA by selecting an siRNA molecule of 19-25 nucleoside bases by selecting a target gene and measuring the functionality of the nucleotide sequences that are complementary to a stretch of nucleotides
comprises a hairpin. The siRNA molecule comprises between 18 and 30 base
                                                                                              Gaps
                                                                                                                                                                                                                                                         ss; siRNA; gene silencing; Bcl-2; optimised; short interfering RNA; RNA interference; firefly; luciferase.
                                                                                              ·:
                                                                           Length 19;
                                                                                                                                                                                                                                                                                                                                                                                                                 ŝ
                                                                                             Indels
                                                                                                                                                                                                                                                                                                                                                                                                                  Stephen
                                                                                                                                                                                                                                         Anti-Firefly luciferase siRNA Luc 77 SEQ ID NO:287.
                                                        BP; 6 A; 3 C; 7 G; 0 T; 3 U; 0 Other;
                                                                           Score 13.8; DB 1;
Pred. No. 9.8e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                 William M,
                                                                                              0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    English.
                                                                                                                                                                                                                                                                                                                                                                                                                  Devin L,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 1; SEQ ID NO 287; 199pp;
                                                                                                                 1203 CCTCTTTCCGGGCTCCA 1219
                                                                                                                                                                                   BP.
                                                                                                                                                                                                                                                                                                                                                                  2002US-0426137P.
2003US-0502050P.
                                                                                                                                                                                                                                                                                                                                                14-NOV-2003; 2003WO-US036787
                                                                            0.8%;
                                                                                    88.2%;
                                                                                                                               19 CGTCTTTCCGTGCTCCA
                                                                                                                                                                                   ADQ60588 standard; RNA; 19
                                                                                                                                                                                                                       (first entry)
                                                                                               Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                  Angela R,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   the target sequence.
                                                                                                                                                                                                                                                                                                                                                                                               (DHAR-) DHARMACON INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2004-420527/39.
                                                                             Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                            WO2004045543-A2
                                                                                                                                                                                                                                                                                                                                                                                                                  Anastasia K,
                                                                                                                                                                                                                                                                                                                                                                   14-NOV-2002;
10-SEP-2003;
                                                                                                                                                                                                                       09-SEP-2004
                                                                                                                                                                                                                                                                                                                              03-JUN-2004.
                                                         Sequence 19
                                                                                              15;
                                                                                                                                                                                                                                                                                        Synthetic.
                                                                                                                                                                                                    ADQ60588;
                                                                                                                                                                 1462
                                                                                               Matches
                                                                                                                                                                RESULT
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Tue Nov

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RESULT 1463

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Human papillomavirus; HPV; amplification; primer;
polymerase chain reaction; PCR; antibody; assay; nitrocellulose filter;
                                                                                                                                                                                                                                                 Avoiding contamination during nucleic acid amplification - using oligo:nucleotide primer contg. unnatural base which can be selectively rendered incapable of further amplification.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Fixing nucleotide sequence to solid support, e.g. nylon filter - using antibody specific for substit. on the sequence as intermediate protein.
                                                                                                                                                                                                                                                                                                                                               Example 1 describes the amplification of HPV-16 DNA by PCR using the primers given in AAQ15430-31 or AAQ15432-33
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The use of probes fixed by antibodies to nitrocellulose filters was
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ·;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                DB 1; Length
                                                                                                                                                                                                                                                                                                                                                                                                Sequence 20 BP; 2 A; 2 C; 7 G; 0 T; 9 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                              Score 13.8; DB 3
Pred. No. 1e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                              0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (INRM ) INSERM INST NAT SANTE & RECH MED,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Disclosure, Page 14; 20pp; French.
                                                                                                                                                                                                                                                                                                                 Example 1; Pag 7; 10pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1308 CAAGACATACAACTACC 1324
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        useful e.g. in pathogen typing.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              BP.
                                                                                                                       90EP-00309492.
                                                                                                                                                    89US-00401840.
                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match 0.8%;
Best Local Similarity 88.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        90FR-00004659
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        90FR-00004659
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        17 CAAGACATACATCGACC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAQ58627 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                15; Conservative
                                                                                                                                                                                    (LIFE-) LIFE TECHN INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Ichen P, Vautherot JF;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (revised)
                                                                                                                                                                                                                    WPI; 1991-067289/10.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 1992-001368/01.
                                                                                                                                                      01-SEP-1989;
                                                                                                                      30-AUG-1990;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            25-MAR-2003
25-APR-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        11-APR-1990;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        11-APR-1990;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           HPV-6 probe
                                                     EP415755-A.
                     Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAQ58627;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          RESULT 1465
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAQ58627
        à
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ;
0
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                                                                                                                                                                                                                                                                                                                                                                                         Human papillomavirus; amplification; primer; polymerase chain reaction;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human papillomavirus; amplification; primer; polymerase chain reaction;
PCR; ss.
siRNA for generating a gene silencing reagent. The present sequence is used in the exemplification of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Avoiding contamination during nucleic acid amplification - using oligo:nucleotide primer contg. unnatural base which can be selectively rendered incapable of further amplification.
                                                                                                                      Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Example 1 describes the amplification of HPV-16 DNA by PCR using the primers given in AAQ15430-31 or AAQ15432-33
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
                                                                                                                      ;
0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      .;
                                                                                  0.8%; Score 13.8; DB 1; Length 19; 88.2%; Pred. No. 9.8e+02; ative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match 0.8%; Score 13.8; DB 1; Length 20; Best Local Similarity 88.2%; Pred. No. 1e+03; Matches 15; Conservative 0; Mismatches 2; Indels
                                                  Sequence 19 BP; 7 A; 3 C; 7 G; 0 T; 2 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Seguence 20 BP; 2 A; 2 C; 7 G; 9 T; 0 U; 0 Other;
                                                                                                                                                 1203 CCTCTTTCCGGGCTCCA 1219
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example 1; Pag 7; 10pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1308 CAAGACATACAACTACC 1324
                                                                                                                                                                                                                                                 :432/c
AAQ15432 standard; RNA; 20 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAQ15430 standard; RNA; 20 BP
                                                                                                                                                                     GTCTTTCCGTGCTCCA 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            90EP-00309492.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         89US-00401840
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CAAGACATACATCGACC 1
                                                                                                                                                                                                                                                                                                                           21-APR-1994 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
                                                                                  Query Match 0.8
Best Local Similarity 88.2
Matches 15; Conservative
                                                                                                                                                                                                                                                                                                                                                           HPV-16 control primer dT1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (LIFE-) LIFE TECHN INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 1991-067289/10.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   HPV-16 primer dUl.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         30-AUG-1990;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         01-SEP-1989;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   21-APR-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          EP415755-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                        Synthetic.
                                                                                                                                                                               17
                                                                                                                                                                                                                                                                                            AAQ15432;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAQ15430
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RESULT 1464

ð 셤 AAQ15430/ 1D AAQ1 XX AC AAQ1 XX DT 21-A XX XX XX HPV-XX KW Huma KW PCR;

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Gaps

Best Loca Matches

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The PCR primer PV3(5') correspond to an oligomer starting at nucleotide 1501 of human papillomavirus type 16. The primer is used to demonstrate a novel in situ PCR method comprising fixed cells, a subset of PCR reagents and opt. a binding protein for single stranded DNA, or fixed cells, a complete set of PCR reagents and the binding protein. The method is used to perform PCR on cells present in histochemical sections or cytochemical smears, e.g. for biological, forensic or pathological studies. The primer was one of a pair used to amplify papillomavirus DNA from human cervical ancer cells silla. A 449 bp PCR prof. was obtd. by this method whereas multiple primer pairs were needed for the same result using conventional pCR methods. See also AAQ34980-6. (Updated on 25-MAR-2003 to correct PN
                                                                                                                                                                                                                                                                                                                                                     Compsn. for in situ polymerase chain reaction on fixed cells - involves preventing reaction until start of thermal cycling, and providing higher sensitivity and selectivity.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          N4-methyl-cytidine; N4-methyl-deoxycytidine; triphosphate; CTP; dCTP; substrate; polymerase; cytosine; oligonucleotide; polynucleotide; sequence analysis; primer extension reaction; PCR; polymerase chain reaction; ampliflication.
                               Amplification; cervical cancer; HPV-16; human papillomavirus;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0.8%; Score 13.8; DB 1; Length 20;
88.2%; Pred. No. 1e+03;
cive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 20 BP; 2 A; 4 C; 6 G; 8 T; 0 U; 0 Other;
                                                                                                                                                                                                                                     (HOFF ) HOFFMANN LA ROCHE & CO AG F. (UYNY ) UNIV NEW YORK STATE RES FOUND.
                                                                                                                                                                                                                                                                                                                                                                                                                               Example 1; Page 10; 14pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1308 CAAGACATACAACTACC 1324
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          BP.
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                                                                                                                                                                                                         91US-00733419.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAQ44798 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (revised)
                                                                                                                                                                                                                                                                                                                          WPI; 1993-028856/04.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               HPV16/pT713 primer.
                                                                                                                                                                                                                                                                                         Bloch W, Nuovo GJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Best Local Similarity
PCR primer PV3(5')
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WO9405684-A1
                                                                                                                                                                       22-JUL-1992;
                                                                                                                                                                                                         23-JUL-1991;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              25-MAR-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  17-MAR-1994.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             29-SEP-1994
                                                                                                                                        27-JAN-1993
                                                                                                    EP524808-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Synthetic.
                                                                    Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAQ44798;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1468
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAQ44798,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RESULT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Oligo:nucleotide-dependent amplification for controlling contamination of prod - by incorporating an exo-sample nucleotide into products.
 exemplified in an assay for HPV. The probes are given in AAO58627-AAO58630 and the primers are given in AAO58631-O58634. (Updated on 25-MAR-2003 to correct PA field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           amplification of a
16) DNA. (Updated on 25-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
                                                                                                                                           Gaps
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88.2%; Pred. No. 1e+03;
tive 0; Mismatches 2; Indels
                                                                                                        Score 13.8; DB 1; Length 20;
Pred. No. 1e+03;
                                                                                                                                         Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                               Polymerase chain reaction; HPV 16; amplification; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The sequence is that of a PCR primer used in the region of the human papilloma virus type 16 (HPV MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 20 BP; 2 A; 2 C; 7 G; 9 T; 0 U; 0 Other;
                                                                                                                                         5;
                                                                      Sequence 20 BP; 5 A; 8 C; 1 G; 6 T; 0 U; 0 Other;
                                                                                                                                         0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                 Human papilloma virus type 16 PCR primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example; Page 10; 18pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1308 CAAGACATACAACTACC 1324
                                                                                                                                                                             1677 CCCCAACTACATCTTCC 1693
                                                                                                                                                                                                           20
                                                                                                                                                                                                                                                                                              BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ВР
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (LIFE-) LIFE TECHNOLOGIES INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CAAGACATACATCGACC 1
                                                                                                        0.8%;
88.2%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        91US-00728874
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                                                                                                                                                                                                4 CCGTAACTACATCTTCC
                                                                                                                                                                                                                                                                                              AAQ34599 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (revised)
(first entry)
                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
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                                                                                                                                           15; Conservative
                                                                                                                                                                                                                                                                                                                                                                 (revised)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity
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                                                                                                        Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       13-JUL-1992;
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26-MAY-1993
                                                                                                                                                                                                                                                                                                                                                               25-MAR-2003
10-MAY-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     EP522884-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       13-JAN-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Hartley JL,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAQ34982;
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Gaps

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deoxycytidine 5'-triphosphare (II) are new. (I) and (II) serve as substrates for RNA and DNA polymerases for incorporation of the N4-(1-4C alky1)-cytosine moiety into oligo- and polymucleotides. They can be used in DNA sequence analysis, primer extension reactions and nucleic acid amplification. To assess the potential for using N4-methyl-dCTP in PCR amplification, reaction mixts. contg. the canonical nucleotide set were compared to mixts. in which dCTP was replaced by the N4-methyl-dCTP in PCR analogue, in a PCR experiment designed to amplify a 293 pp sequence of HPV16 DNA. Using a high-temp. regimen the desired fragment was obtained with the canonical dNTPs, but not with N4-methyl-dCTP, A low-temp. regimen, conducted with dCTP or with N4-methyl-dCTP, A low-temp. regimen, conducted with dCTP or with N4-methyl-dCTP, A low-temp. mixt., cleanly produced identical amts. of the expected fragment as the sole amplification product. (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Cpds. N4-(1-4C alkyl) cytidine 5'-triphosphate (1) and N4-(1-4C alkyl)-2'
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Peptide transport gene, atptr2a, disease-resistance, fungus-resistance, insect-resistance, pathogen-resistance, herbicide-resistance, transgenic plant, crop improvement, polymerase chain reaction, primer,
                                                                                                                                                                                                                                                                                                                                                New N4-alkyl-(deoxy) cytidine 5'-tri:phosphate cpds. - useful in sequence analysis, primer extension reactions and nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0.8%; Score 13.8; DB 1; Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 20 BP; 2 A; 2 C; 7 G; 9 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Peptide transport gene atptr2a PCR primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Disclosure; Page 12; 40pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1308 CAAGACATACAACTACC 1324
                                                                                                                                     (LIFE-) LIFE TECHNOLOGIES INC
93WO-US008145.
                                                                 92US-00941370.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     88.28;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CAAGACATACATCGACC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Arabidopsis thaliana
                                                                                                                                                                                                                                                                               WPI; 1994-101109/12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 1995-336935/43
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Local Similarity
les 15; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                    amplification.
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30-AUG-1993;
                                                                 04-SEP-1992;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 RT-PCR; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAT05336
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XX AATO
XX AATO
XX XX AATO
XX PEPT
XW FEPT
XW TT-E
XW TT-E
XX ATA
XX ATA
PN WO9!
XX YX PO 21-:
XX YX PO 21-:
XX PD 21-:
XX
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2; Indels

Pred. No. 1e+03; 0; Mismatches

ВР

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An upstream primer (AAT05336) starting at base 1975 of the Arabidopsis thaliana peptide transport atptr2a gene (see AAT05334) and a downstream primer (AAT05337) starting at base 2528 were used in RT-PCR to measure the extent of ATPTR2A transcription in plant tissue. A 569 bp fragment of the atptr2a open reading frame is generated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Pigment epithelium-derived factor (PEDF) has both neuronotrophic and gliastatic activity, making it useful in cases where neurons die quickly and gliastatic activity, making it useful in cases where neurons die quickly neuroglia tend to proliferate (gliosis), e.g. in CNS cell culture, in neurodegenerative diseases and in CNS injury. The neuronotrophic effect of PEDF is especially useful for enhancing survival of neuronal cell of PEDF is especially useful for enhancing survival of neuronal cell human foetal brain cells and neural retina and photoreceptor cells. The gliastatic activity of PEDF can be applied to inhibiting glial cell proliferation in certain tumours. Antibodies directed against PEDF can be used for inhibiting PEDF activity or in an immunoassay for determining levels of PEDF in fluid, cellular or tissue samples e.g for determining ageing and/or other degenerative diseases. Eight primers (AAT11661-68)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           88
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Use of pigment epithelium derived factor - for enhancing neuronal cell survival and inhibiting glial cell proliferation, useful, e.g. in CNS cell culture or to treat neuro-degenerative diseases.
Plant peptide transport genes - used to increase plant resistance to herbicidal peptide(s), pref. those produced by a plant pathogen.
                                                                                                                                                                                                                                                                                                    Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Pigment epithelium-derived factor; PEDF; neuronal cells; neurons; glial cells; gliastatic; gliosis; central nervous system; CNS; neurodegenerative disease; injury; neuronotrophic; brain cells; Parkinson's disease; photoreceptor cells; retina; inhibition; proliferation; immunoassay; antibody; ageing; degenerative disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Primer for amplifying pigment epithelium-derived factor fragment.
                                                                                                                                                                                                                                                                                                  .
0
                                                                                                                                                                                                                                                         0.8%; Score 13.8; DB 1; Length 20;
18.2%; Pred. No. 1e+03;
ve 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Taniwaki T;
                                                                                                                                                                                                                   Sequence 20 BP; 3 A; 6 C; 4 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (USSH ) US DEPT HEALTH & HUMAN SERVICES.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example 8; Page 38; 151pp; English.
                                                          Example 8; Page 40; 79pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 BP
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94US-00367841
                                                                                                                                                                                                                                                                             88.2%;
                                                                                                                                                                                                                                                                                                                                        6 GCAGCGTAAAGGATGGA
                                                                                                                                                                                                                                                                                                                                                                               GCAGCGTAAATCATGGA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAT11661 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
                                                                                                                                                                                                                                                  Query Match
Best Local Similarity 88.27
Matches 15, Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Chader GJ, Becerra SP,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 1996-039966/04.
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30-DEC-1994;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             16-JAN-1997
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAT11661;
                                                                                                                                                                                                                                                                                                                                                                                                                                       RESULT 1470
                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAT11661
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0
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for H chain of Fas specific antibody coding sequence.

(first entry)

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This sequence represents a primer for the coding sequence for the protein of the invention is a recombinant protein (A), that comprises at least one region corresponding to an immunoglobulin (Ig) variable region which enables the protein to recognise and specifically bind to an artigen, preferably human Fas, and has substantially no more immunogenicity in a human patient than a human antibody. The proteins are useful for treating autoimmune diseases, especially rheumatism (rheumatoid arthritis). (A) is based on a murine monoclonal antibody. As the protein lacks the constant region, it has autibody antibody.
                                                                                       Fas; antibody; human; immunoglobulin; variable region; rheumatism; autoimmune disease; rheumatoid arthritis; therapy; CDR; heavy chain; complementarity determining region; PCR primer; amplify; ss.
                                                                                                                                                                                                                                                                                                                                                                         Anti-Fas recombinant antibodies - useful fidiseases, especially rheumatoid arthritis.
                                                                                                                                                                                                                                                                                                                                                                                                                  Example 4; Page 16; 72pp; English.
                                                                                                                                                                                                                                                                                                                        Serizawa N, Ichikawa
                                                                                                                                                                                                                                                                                              (SANY ) SANKYO CO LTD
                                                                                                                                                                                                                                                                                                                                                WPI; 1997-482673/45.
                                                                  Primer SHR-16
                                                                                                                                                                                                                                          27-MAR-1997;
                                                                                                                                                                                                                                                                   01-APR-1996;
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                                        15-APR-1998
                                                                                                                                                                                     EP799891-A1
                                                                                                                                                                                                                08-OCT-1997.
                                                                                                                                                Synthetic.
             AAV03721;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAT47350;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
                                                                                                                                                              ds snw
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RESULT 1473
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Neo-specific primer P8, (AAT78982), primer P586 (AAT78983) derived from exon 5 of the mouse Huntington's disease (HD) gene (see AAT78974), and primer P9 (AAT78984) derived from intron 5 of the gene were used in the genotype analysis of heterozygous transgenic mice embryos carrying a targeted mutation in exon 5. The results indicated that loss of function of the endogenous Hdh gene resulted in embryonic lethality during early post-implantation development. Transgenic mice can be used as models of
were synthesised base on the CDNA sequence of PEDF and used to amplify fragments of the PEDF gene for later sequencing. Two primers (AAT11661, AAT11662) were used to amplify a 2 kilobase fragment from exon 3 to exon
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mouse Huntington's Disease gene - useful for generating transgenic mice as a model of Huntington's Disease.
                                                                                                                                                                                                                                                                                                                                                  ington's disease; animal model; transgenic animal; mouse; therapy; screening; mhd gene; polymerase chain reaction; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
                                                                                                                        Gaps
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0
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                                                                                          Score 13.8; DB 1; Length 20; Pred. No. 1e+03; 0; Mismatches 2; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Indels
                                                                                                                                                                                                                                                                                                                        Mouse Huntington's disease gene exon 5 primer P586.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                5;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 20 BP; 1 A; 5 C; 7 G; 7 T; 0 U; 0 Other;
                                                                  G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1666 CACAGGGCAGCCCCCAA 1682
                                                                                                                                                 1631 CCAGCAGGCAGCGCTG 1647
                                                                                                                                                                                                                                            BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (UYBR-) UNIV BRITISH COLUMBIA.
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                                                                    ,
                                                                                             Match 0.8%;
Local Similarity 88.2%;
les 15; Conservative
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                                                                     BP; 4 A; 6 C;
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Lin B,
                                                                                                                                                                                                                                                                                                                                                    Huntington's
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        03-JUN-1996;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 15;
                                                                      Sequence 20
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                                                                                                                                                                                                                                                                                                                                                                                            Synthetic.
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                                           of PEDF
                                                                                                                                                                                                                                                                   AAT78983
                                                                                                 Query Match
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for treating auto-immune

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Yonehara

Nakahara K,

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97EP-00302415. 96JP-00078570.

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                                                                                      Gaps
                                                                                      0
                                       Score 13.8; DB 1; Length 20;
Pred. No. 1e+03;
                                                                                      Indels
                                                                                                                                                                                                                                                                                                                                                                                                                            Variant #6 of universal primer sequence for M13mp18.
Sequence 20 BP; 2 A; 8 C; 2 G; 8 T; 0 U; 0 Other;
                                                                                   5;
                                                                                      0; Mismatches
                                                                                                                                  1452 TCCATTCTTCCTCAGTC 1468
                                                                                                                                                                            20
                                                                                                                                                                                                                                                                                         BP.
                                         0.8%;
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                                                                                        15; Conservative
                                                                 Local Similarity
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AAV03721 standard; DNA; 20

RESULT 1472 AAV03721 ID AAV0372

20

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(HOFF ) HOFFMANN LA ROCHE & CO AG F.
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                                                                                                                                                                          psoriasis.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAT47345-T47374 represent variants of a universal primer sequence (see AAT47344) derived from the bacteriophage vector M13mp18. This sequence can be used as half of the DNA primer of the invention. The primers are used for amplification of a target DNA sequence, and can be used in a multiplex PCR amplification. The primers have the sequence 5'-Xy-3', where X is a sequence that does not hybridise to the target sequence (such as this sequence. The melting temperature of a hybrid between X and the target sequence. The melting temperature of a hybrid between X and the target sequence. The melting temperature of a hybrid between X and the target sequence. The melting temperature of a hybrid between X and the target sequence. The melting temperature of a hybrid between X and the complement (in the absence of other sequences) is 60 degrees C. During early cycles of amplification, products are synthesised that contain the chimeric primers on either end. The primers then serve as high stringency recognition sequences for subsequent rounds of thereby reducing preferential amplification of certain targets. The containing Artification of any integer specific domain. They are used for the simultaneous PCR amplification of multiple Odmain. They are used for the simultaneous PCR amplification of multiple community as particularly useful in high-throughput genetic screening for detecting the presence of multiple defined targets e.g. to detect mutations in genes like the cystic fibrosis transmembrane conductance regulator (CPTR), the Wilms Tumour, and the beta-thalassaemia genes
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0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Puromycin-sensitive aminopeptidase, PSA, treatment, cancer, psoriasis, proliferative disorder, hybridise, antisense oligonucleotide, ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Puromycin-sensitive aminopeptidase (PSA) antisense oligonucleotide 2.
                                                                                                                                                                                  Universal primer used for multiplex DNA amplification - allows simultaneous amplification of multiple DNA target sequences for high through-put genetic screening.
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88.2%; Pred. No. 1e+03;
ative 0; Mismatches
                                                                                                                                                                                                                                         Claim 8; Page 10; 38pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1182 TGAGATGGCCACAGGCC 1198
                                                    96WO-US009637.
                                                                            95US-00474450
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TGACATGGCCACCGGCC 3
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                                                                                                      (GENZ ) GENZYME CORP
                                                                                                                                                           WPI; 1997-052372/05.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity
Matches 15; Conserv
                                                                             07-JUN-1995;
                                                    06-JUN-1996;
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Homo sapiens
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                          19-DEC-1996
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This antisense oligonucleotide is specifically hybridisable with selected DNA or RNA deriving from the puromycin-sensitive aminopeptidase (PSA)-99. This oligonucleotide is used for dignosing conditions associated with PSA expression. The human PSA-99 (875 amino acids) and the murine PSA-99 (920 amino acids) both exhibit PSA activity and can be used to generate anti-CSC amino acids) both exhibit PSA activity and can be used to generate anti-CSC amino acids) both exhibit PSA activity and can be used to generate anti-CSC amino acids) both exhibit PSA activity or departs which act by targeting PSA, for modulating PSA activity or cunction. They can be used to influence proteolytic degradation of endogenous PSA substrates, proliferation rate or viability of cells or to induce apoptosis within cells by inhibiting PSA activity. Agents which can diminish PSA activity in cells, by modulation of the amount of PSA synthesis, may be used in treatment of proliferative diseases, including tumours such as leukaemias and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ö
                                                                                                                                                                                                                                                                                                                                                                               Isolated protein with puromycin-sensitive aminopeptidase activity - which may be used in treatment of proliferative disorders, including cancer and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
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88.2%; Pred. No. 1e+03;
ve 0; Mismatches 2; Indels
                                                                                                                                                                                                                                 Altmann K, Schlapbach
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     carcinomas or epithelial disorders like psoriasis
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                                                                                                                                                                                                                                 Tobler AR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 36; Page 109; 141pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 179 GAGGCATAGACAAGACC 195
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96WO-EP001518.
                                                                           96WO-EP001518.
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nes 15; Conservative
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                                                                                                                                                      (NOVS ) NOVARTIS AG.
                                                                                                                                                                                                                                                                                                    WPI; 1997-512727/47
09-APR-1996;
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Matches
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NAME OF THE PROPERTY OF THE PR
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                                                                                                                                                                                                                                         Sequences shown in AAV33257 to AAV3263 represent primers used for the PCR amplification of the human papillomavirus (HPV) type 16 genome contained in the human cervical cancer cell line SiHa. The invention provides a thermal cycler sample compartment optimised for holding and provides a thermal cycler sample compartment optimised for holding and provides a thermal cycler sample compartment optimised for holding and controlling the temperature of one or more microscopes which facilitates a device (barrier) for protecting a microscope slide from fluid or vapour when the slide is scaled in the device, comprising a plastics material that has high thermal conductivity, and is impervious to fluid or vapour, and is dimensioned so as to receive the slide. The new thermal cycling compartment is useful for performing in situ PCR for detection of target nucleic acid sequences directly from cells fixed onto a microscope slide, used in the field of cell biology, forensic science and clinical, veterinary and plant pathology. The modified heat blocks increase the speed and reliability of in situ PCR performed on microscope slides by accelerating and rendering more uniform the heat transfer which occurs during thermal cycling.

(Updated on 25-MAR-2003 to correct PP field.) (Updated on 25-MAR-2003 to
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LRP5; LDL-receptor related protein; LRP-3; IDDM; diagnosis; endocytosis; insulin dependent diabetes mellitus; autoimmune disease; glomerulonephritis; inflammation; viral infection; osteoporosis; hypercholesterolemia; Alzheimer's disease; low density lipoprotein;
                                                                                                                                    New thermal cycler for in-situ PCR on microscope slides - and device for protecting microscope slides from fluid or vapour.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Hammond H;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 13.8; DB 1; Length 20;
Pred. No. 1e+03;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 20 BP; 2 A; 4 C; 6 G; 8 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mouse LRP-3 cDNA PCR primer 378r (mulrp3 3r).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COX RD,
  (UYNY ) UNIV NEW YORK STATE RES FOUND.
                                                                                                                                                                                                        Example 1; Page 10; 16pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1308 CAAGACATACAACTACC 1324
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Caskey CT,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             97US-0043553P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ო
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                98WO-GB001102
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (WELL ) WELLCOME TRUST LID.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  19 CAAGACATACATCGACC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAV85967 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (MERI ) MERCK & CO INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Hess JW,
                                               Nuovo GJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          correct PR field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PCR primer; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  15-APR-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             15-APR-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       05-JUN-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WO9846743-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              10-FEB-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   22-OCT-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               rodd JA,
                                            Bloch W,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAV85967;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mus ap
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       RESULT 1476
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      g
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The present invention describes LRP5 (low density lipoprotein (LDL) receptor related protein, previously designated LRP-3). Nucleic acid molecules (NAMS) encoding LRP5 can be used for determining if an individual is susceptible to insulin dependent diabetes mellitus (IDDM). The NAMS or proteins can be used for reducing triglyceride levels in the serum of an individual. Therapies that affect LRP5 may also be useful in the treatment of autoimmune diseases such as glomerulonephritis, diseases and disorders involving disruption of endocytosis and/or antigen presentation, cytokine clearance and/or inflammation, viral infection, pathogenic bacterial toxin contamination, elevation of free fatty acids or hypercholesterolemia, type 2 diabetes, osteoporosis, Alzheimer's disease and cardiovascular disease. Products from the present invention can also be used for detection, diagnosis, and drug screening. AAVBS917 to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                can also be used for detection, diagnosis and drug screening. AAV85917 to AAV86012 represent PCR primers for obtaining LRP-3 CDNA
                                                                                                                                              New isolated LDL-receptor related protein - used to develop products for treating, e.g. elevated triglyceride levels, diabetes, autoimmune disorders, inflammation or Alzheimer's disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               changes in mRNA cancer-associated gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The primers AAV43732-V43776 were to produce cancer associated gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ss; cancer; PCR; Northern blotting; ribonuclease protection assay; diagnosis; metastatic cancer; primer; amplification.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ;
0
Nakagawa Y;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0.8%; Score 13.8; DB 1; Length 20; 88.2%; Pred. No. 1e+03; Live 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Kato I;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ples - by
specific
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 20 BP; 1 A; 6 C; 5 G; 8 T; 0 U; 0 Other;
   Metzker ML,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Detection of cancer cells in tissue samples expression compared to normal tissue of spec
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Œ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Hino
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Disclosure; Page 67; 92pp; Japanese
   Merriman TR,
                                                                                                                                                                                                                                                                                     Claim 12; Page 117; 200pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Α,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Cancer associated gene primer 2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1435 GAGGATGCCATGAACA 1451
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Asada
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            98WO-JP000667.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      97JP-00052508
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (TAKI ) TAKARA SHUZO CO LID.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   20 GAGGAGGCCATCAAACA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAV43733 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
                             Twells RCJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mukai H,
Kawaguchi Y,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 1998-467552/40.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity
                                Phillips MS,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Yoshikawa Y,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               16-NOV-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WO9837187-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            18-FEB-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         21-FEB-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  27-AUG-1998.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         sequences.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAV43733;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    RESULT 1477
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Gaps

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Query Match 0.8%; Score 13.8; DB 1; Length 20; Best Local Similarity 88.2%; Pred. No. 1e+03; Matches 15; Conservative 0; Mismatches 2; Indels

1308 CAAGACATACAACTACC 1324

m

19 CAAGACATACATCGACC

d

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Sequence 20 BP; 2 A; 4 C; 6 G; 8 T; 0 U; 0 Other;

microscope slide

SXS

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0
fragments which can be used to detect cancer cells in tissue samples or biological fluids. They are detected by monitoring the change in mRNA expression as compared to normal tissue of one or more cancer-associated genes whose cDNA stringently hybridises to the nucleic acid fragments. The change in expression may be an increase or a decrease compared to normal tissue. The mRNA expression may be determined by PCR, Northern blotting or ribonuclease protection assay, or by determining the change in the amount of protein encoded by the gene(s) as compared to normal tissue, for example by using a labelled antibody recognising the protein. Detection of cancer cells for cancer diagnosis, including detection of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human papillomavirus; HPV; thermal cycling device; ceramic sample plate; biological sample; thermal sensor; heater; cooler; thermal cycling; rapid heat transfer; microscope slide; PCR amplification; hybridisation; target nucleic acid; PCR primer; ss.
                                                                                                                                                  metastatic cancer cells in tissues other than the primary tumour site
                                                                                                                                                                                                                                          Gaps
                                                                                                                                                                                                                                          ..
                                                                                                                                                                                                          0.8%; Score 13.8; DB 1; Length 20;
88.2%; Pred. No. 1e+03;
trive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human papillomavirus (HPV) gene amplifying primer PV3.
                                                                                                                                                                               Sequence 20 BP; 2 A; 2 C; 7 G; 9 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                   1055 AGTCAATCCCAACAAAG 1071
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 97US-00810641.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      98WO-US004041.
                                                                                                                                                                                                                                                                                                 17 AGTCAACCCAAACAAG 1
                                                                                                                                                                                                                                                                                                                                                                       AAV54679 standard; DNA; 20
                                                                                                                                                                                                Ouery Match
Best Local Similarity 88.2%
                                                                                                                                                                                                                                                                                                                                                                                                                                 13-NOV-1998 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Chaplin J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (MINU ) UNIV MINNESOTA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human papillomavirus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 1998-495869/42.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Blumenfeld M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   03-MAR-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      03-MAR-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WO9839479-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         11-SEP-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                    AAV54679;
                                                                                                                                                                                                                                                                                                                                            RESULT 1478
                                                                                                                                                                                                                                                                                                                                                        8X3333333333X8
                                                                                                                                                                                                                                                                                                셤
                                                                                                                                                                                                                                                                      8
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Human, c-fos; c-jun, activating protein 1; AP-1; diagnosis, metastasis; antisense oligonucleotide; phosphorothioate; regulation; malignant tumour; cell cycle expression; hyperproliferative disease; ss.

Human c-jun protein antisense oligonucleotide #9.

(first entry)

04-FEB-1999

AAV69985;

BP

AAV69985 standard; DNA; 20

RESULT 1479

AAV69985

/*tag= a /note= "phosphorothioate linkages"

98WO-US007386.

14-APR-1998;

Location/Qualifiers

modified base

Key

WO9846272-A1

22-0CT-1998

Homo sapiens

Synthetic

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AAV69978 to AAV69988 represent antisense oligonucleotides which are specifically hybridisable with a region of a nucleic acid encoding human c-Jun protein. The antisense compound regulates the expression of the c-Jun protein. The present invention also describes antisense oligonucleotides which regulate the c-Fos protein. The antisense oligonucleotides are used for the diagnosis and treatment of diseases or disorders associated with Activating Protein I expression, of which c-Fos and c-Jun are subunits. The antisense oligonucleotides are used in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Antisense oligonucleotides regulating Activating Protein 1 subunits - hybridise with c-fos and c-jun mRNA, used for regulating metastasis, cell cycle expression and hyperproliferative disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      chemotherapeutic agent. They are used to regulate the expression of c-Fos or c-Jun in cells or tissues, preferably by inhibiting metastasis. They also regulate cell cycle expression and can be used to treat an animal with, or being prone to, a hyperproliferative disease
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 compositions as c-Fos and/or c-Jun together with a carrier and a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DB 1; Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 20 BP; 2 A; 11 C; 5 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0.8%; Score 13.8; DB;
88.2%; Pred. No. 1e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Baker B;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 12; Page 71; 120pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Dean NM, Mckay R, Miraglia L,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 97US-00837201.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 1998-609906/51.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 14-APR-1997;
PCR amplification of the Human papillomavirus (HPV) gene contained in the human cervical cancer cell line SiHa. These are used in the course of the invention which provides a thermal cycling device comprising a ceramic sample plate. This device has a ceramic sample plate supporting a flat substrate carrying a biological sample and a thermal sensor, a heater thermally coupled to the plate and a cooler for the substrate. The device cither maintains the temperature of the sample or subjects it to thermal cycling. The thin ceramic plate permits very rapid heat transfer to a sample on a microscope slide and this thermal cycling devise can be used for PCR amplification or hybridisation of target nucleic acid on
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Thermal device for PCR amplification or hybridisation of target nucleic acid on microscope slide - has ceramic sample plate supporting flat substrate for sample and heater and cooler controlled to maintain or rapidly cycle temperature of sample.

Sequences shown in AAV54677 to AAV54683 represent primers used for the

Example 2; Page 34; 58pp; English.

transgenic; transgenic food; ss.

Simons LH;

Goddijn OJM, 1, Elzinga N;

à g

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The invention relates to a vector for plant transformation, comprising a T-DNA with flanking T-DNA borders and also a polynucleotide that prevents the development of plant transformants containing more vector sequences than the T-DNA sequence. The vectors encode toxins that are specific to certain plants, or those which act on membrane systems and /or other cellular structures. Examples of genes include those encoding ribozymes against endogenous RNA transcripts, proteins evoking hypersensitive reactions, and RNA transcripts used for antisense/co- suppression inhibition of gene expression. The polynucleotide sequence contained in the vectors prevents the transfer of DNA sequences beyond the T-DNA borders. This avoids contamination of transgenic plants and/ or transgenic food with vector DNA. Sequences AAX05690-91 represent primers used for the PCR amplification of the barnase open reading frame. This is
                                                                                                                                                                                                                                                                                                                           New vector for plant transformation - useful for producing toxins that are specific to certain plants, or those which act on membrane systems and/or other cellular structures.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                HIV cofactor inhibitor; HIV infection; CXCR4 gene; CCR5 gene; drug composition; antisense; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           used in the construction of a barnase expression cassette
Barnase open reading frame fragment amplifying primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Seguence 20 BP; 5 A; 5 C; 5 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 13.8; DB 1;
Pred. No. 1e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CCR5 gene inhibiting antisense oligo AS(s)-60.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0; Mismatches
                                                                                                                                                                                                                                                   Ohl SA, Go
Tigelaar H,
                                 toxin;
                                            binary vector; PCR primer; barnase;
                                                                                                                                                                                                                                                                                                                                                                                     Example 4; Page 21; 34pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              131
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 BP.
                                                                                                                                                                98WO-EP004171.
                                Plant transformation; T-DNA;
                                                                                                                                                                                            97EP-00201990
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88.2%;
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                                                                                                                                                                                                                                                    Ponstein AS,
Hoekstra S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
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                                                                                                                                                                                                                         (MOGE-) MOGEN INT NV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Local Similarity
                                                                                                                                                                                                                                                                                               WPI; 1999-106063/09.
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                                                                                                        WO9901563-A1
                                                                                                                                                                29-JUN-1998;
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                                                                                                                                     14-JAN-1999
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                                                                                                                                                                                                                                                    Stuiver MH,
                                                                                                                                                                                                                                                                 Dekker BMM
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                                                                            Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
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    o.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   e.g. to
or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gene-specific primer PK22L234 and adapter primer API (see AAV32935) were used in a RACE FCR technique for cloning a 5' coding region of novel human cyclin-dependent kinase 10 (CDK10) CDNA, using adapter-ligated human placenta cDNA as template. Nested primers (see AAV32956-37) were used in a second PCR, to produce an approximately 600 bp product. A 3' fragment was identified by database search, and a full-length sequence (see AAV32932) was produced in vector pLITMUS28:CDK10. The CDK10 protein product (see AAV3893) is used e.g. to identify modulators of cell cycle progression and for treating cancer or immune cell proliferation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
      Gaps
                                                                                                                                                                                                                                                CDK10; cyclin-dependent protein kinase; cell cycle; human; cancer; cell proliferation; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New nucleic acid encoding human cyclin-dependent kinase-10 - used identify modulators of cell cycle progression for treating cancer immune cell proliferation.
                                                                                                                                                                                                                         Human cyclin-dependent protein kinase CDK10 cDNA primer PK22L234.
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Pred. No. 1e+03;
0; Mismatches 2; Indels
    Indels
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      5,
      Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Example 1; Page 27; 58pp; English.
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                                                                                                                                     AAV32934 standard; DNA; 20 BP.
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                                 552 GCCCCTCAGCCGCCGCC
                                                             2 GCCCCTCAGCCCCCGAC
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      Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                              & CO INC.
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Best Local Similarity
Matches 15; Conserv
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AAX05691 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (MERI ) MERCK
                                                                                                                                                                                                                                                                                                               Homo sapiens.
                                                                                                                                                                                                                                                                                                                                             WO9835015-A1
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      15;
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                                                                                                                                                                                                                                                                                                  Synthetic.
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                                                                                                           RESULT 1480
      Matches
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Indels

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Length 20;

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RESULT 1484
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                                                                                                                                                oligonucleotides with a base sequence complementary to the CXCR4 or CCR5 genes. Such inhibitors can be formulated into drug compositions for prevention or treatment of HIV infection, with inhibition of expression of CXCR4 or/and CCR5 gene. Sequences AAZ31244-306 represent antisense
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PCR primers AAZ01426-Z06209 were used to amplify open reading frames (ORFs) of the genome of Chlamydia trachomatis (see AAZ01425). These ORFs
                                                                                                                                                                                                                                            Gaps
                                                                                          Antisense oligonucleotide-based HIV cofactor inhibitors, as drug compositions for treatment of HIV infection.
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                                                                                                                                     The invention provides HIV cofactor inhibitors that contain
                                                                                                                                                                                                                          1; Length 20;
                                                                                                                                                                                                                                                                                                                                                                                    PCR primer used to amplify an ORF of Chlamydia trachomatis.
                                                                                                                                                                                                                                            Indels
                                                      Wada A;
                                                                                                                                                                                                       Seguence 20 BP; 3 A; 6 C; 6 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                         Score 13.8; DB 1
Pred. No. 1e+03;
                                                      Takai K,
                                                                                                                                                                                                                                           Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 sequence of Chlamydia trachomatis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Disclosure, Page 1671; 1755pp; English.
                                                      Kimura T,
                                                                                                                                                                           of CXCR4 or/and CCR5 gene. Sequence oligonucleotides to the CCR5 gene
                                                                                                                     Claim 6; Page 16; 59pp; Japanese
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                                                                                                                                                                                                                                                              92 CTGAGGTTGCTCGCGCG 108
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98US-0107077P.
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99WO-JP001722
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                  98JP-00125452
                                                                                                                                                                                                                                                                                                                             AAZ04231 standard; DNA; 20
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                                    (MARI -) MARINE BIO CO LTD
                                                                                                                                                                                                                                            Conservative
                                                      Takaku H, Yamamoto N,
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Synthetic.
Chlamydia trachomatis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 1999-371125/31.
                                                                        WPI; 1999-620207/53.
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Best Local Similarity
Matches 15; Conserv
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04-NOV-1998;
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                                                                                                                                                                                                                                                                                                                                                AAZ04231;
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against Chlamydia trachomatis. Antisense and ribozyme sequences can also be used to control growth of the microcranism. Chlamydia trachomatis is increased to control growth of the microcranism. Chlamydia trachomatis is responsible for a large number of diseases, e.g. eye diseases such as conventional trachoma, nonendemic trachoma, paratrachoma, and inclusion epidymicis; genital diseases such as nongonococcal uretritis, epidial diseases such as nongonococcal uretritis, premumopathy in breast feeding infants; and venereal lymphogranulomatosis. The polypeptides of the invention may be of use in treating these diseases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Vaccine, eye disease; conventional trachoma; nonendemic trachoma; paratrachoma; inclusion conjunctivitis; genital disease; perihepatitis; nongonococcal uretritis; epidymitis; cervicitis; salpingitis; PCR primer; bartholinitis; pneumopathy; venereal lymphogramulomatosis; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PCR primers AAZ01426-Z06209 were used to amplify open reading frames (ORFs) of the genome of Chlamydia trachomatis (see AAZ01455). These ORFs encode polypeptides (see AAY36754-Y37949) which can be used as vaccines against Chlamydia trachomatis. Antisense and ribozyme sequences can also be used to control growth of the microorganism. Chlamydia trachomatis is responsible for a large number of diseases, e.g. eye diseases such as conventional trachoma, nonendemic trachoma, paratrachoma, and inclusion conjunctivitis; genital diseases such as nongonococcal uretritis, originatis, cervicitis, allpingitis, perihopathicis, barbingitis; peneumopathy in breast feeding infants; and venereal lymphogranulomatosis. The polypeptides of the invention may be of use in treating these
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 20 BP; 4 A; 6 C; 4 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0.8%; Score 13.8; DB 1
88.2%; Pred. No. 1e+03;
iive 0; Mismatches
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97FR-00016034.
98US-0107077P.
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Best Local Similarity 88.2
Matches 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Chlamydia trachomatis.
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17-DEC-1997;
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Deletion sequence oligonucleotide; sensor array; eukaryotic pathogen; probe; cellular adhesion modulator; cellular proliferation modulator; human retrovirus; human immunodeficiency virus; non-human retrovirus;
                                                                                                                          Deletion sequence oligonucleotide 2.
                            AAX23549 standard; DNA; 20
                                                                                           (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                       (ISIS-) ISIS PHARM INC.
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                                                                                                                                                                                                             HIV; primer; ss
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                                                                                             18-JUN-1999
                                                                                                                                                                                                                                          Synthetic.
                                                             AAX23549;
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                                                                                                                                                                                                                                                                                                                                                                                                                                          Chen D,
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             AAX23549,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PCR primers AAZ01426-Z06209 were used to amplify open reading frames code polypeptides (see AAY36754-Y37949) which can be used as vaccines against Chlamydia trachomatis, which can be used as vaccines against Chlamydia trachomatis. Antisense and ribozyme sequences can also be used to control growth of the microorganism. Chlamydia trachomatis is responsible for a large number of diseases, e.g. eye diseases such as conventional trachoma, nonendemic trachoma, paratrachoma, and inclusion conjunctivitis; genital diseases such as nongoncoccal uretritis, practicitis, cervicities, salpinglitis, perihepatitis, bartholinitis; pneumopathy in breast feeding infants; and venereal lymphogramulomatosis.
                                                                                                                                                                                                                                                                                                                                        Vaccine; eye disease; conventional trachoma; nonendemic trachoma; paratrachoma; inclusion conjunctivitas; genital disease; perihepatitis; nongonococcal uretritis; epidymitis; cervicitis; salpingitis; PCR primer; bartholinitis; pneumogathy; venereal lymphogranulomatosis; ss.
                                                                Gaps
                                                              ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0.8%; Score 13.8; DB 1; Length 20;
                                                                                                                                                                                                                                                                                                         PCR primer used to amplify an ORF of Chlamydia trachomatis.
                            0.8%; Score 13.8; DB 1; Length 20; 88.2%; Pred. No. le+03;
                                                             Indels
                                                             2;
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BP; 7 A; 3 C; 7 G; 3 T; 0 U; 0 Other;
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; Mismatches
                                                              0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Disclosure; Page 1754; 1755pp; English.
                                                                                               1433 CAGAGGATGCCATGAAA 1449
                                                                                                                 CAGAGGATGCTGTGAAA 17
                                                                                                                                                                                                               BP.
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97FR-00016034.
98US-0107077P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        98WO-IB001939
                                                                                                                                                                                                             AAZ05240 standard; DNA; 20
                                                                                                                                                                                                                                                                            (first entry)
                                                                15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                            Chlamydia trachomatis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 1999-371125/31.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Best Local Similarity
Matches 15; Conser
                                             Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (GEST ) GENSET
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WO9928475-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           27-NOV-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          28-NOV-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        04-NOV-1998;
Sequence 20
                                                                                                                                                                                                                                                                            07-0CT-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           10-JUN-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           17-DEC-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Griffais R;
                                                                                                                                                                                                                                                                                                                                                                                                                             Synthetic.
                                                                                                                                                                                                                                              AAZ05240;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
                                Query Match
                                                                                                                                                                                RESULT 1485
                                                                Matches
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98WO-US018084.

97US-00923771

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This invention describes a novel composition comprising a number of
sensor arrays, where each array comprises a unique probe oligonucleotide,
which is the reverse complement of part of a unique target
which is the reverse complement of part of a unique probe oligonucleotide probe oligonucleotides of the arget deletion sequence
oligonucleotides The compositions form a method for characterizing a
sample of target deletion oligonucleotides of the sensor arrays. Such
hybridize with the probe oligonucleotides of the sensor arrays. Such
oligonucleotides characterized by the method form pharmaceutical
compositions that are useful for modulating cellular adhesion or
proliferation, and being active against a eukaryotic pathogen, a human immunodeficiency virus (HIV), or a non-human
retrovirus, a human immunodeficiency virus (HIV), or a non-human
retrovirus, including influenza virus, Epstein-Barr virus, Respiratory
Syncytial Virus or cytomegalovirus (CMV). The compositions enable
characterization of deletion sequences, and quantification of different
species of deletion sequences, and quantification of an anixture
species of deletion sequences, and plantification of an anixture.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           .;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              reverse complement is not modified, the method may be performed using
                                                                  New compositions comprising sensor arrays made up of unique probe oligonucleotides - useful for characterizing a sample of target deletion
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Also, if the specificity of the oligonucleotide's nucleobase sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ·,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0.8%; Score 13.8; DB 1; Length 20; 18.2%; Pred. No. 1e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 20 BP; 0 A; 5 C; 5 G; 10 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mismatches
                                                                                                                                                                                                                          Example 1; Page 89; 163pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         133 ATGAAGAAGATCAAACG 149
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAX92036 standard; DNA; 20 BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        88.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAGAAGAAGAGCAAACG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             oligodeoxynucleotides
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Local Similarity
WPI; 1999-205198/17.
                                                                                                                                                    oligonucleotides.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         RESULT 1487
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ID AAX9
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The invention provides a DNA molecule comprising a sequence of nucleotides encoding an alphalF-subunit of a mammalian retinal calcium channel (RCC), including a human abhalF-subunit, a murine alphalF-subunit and orthologs of the human and murine alphalF-subunits. The RCC gene can orthologs of the human and murine alphalF-subunits. The RCC gene may be used to develop products for diagnostic tests, for incomplete CSMB and risk assessment in affected families. The RCC gene can provide information as to the basic defect in this retinal conditions, which could lead to effective methods for treatment or cure of the disorder. As the associated features of myopia, nystagmus and strabismus frequently observed in patients with incomplete CSMB may be caused by calciumregulated development pathways, identification of the RCC gene may help to elucidate the molecular details of eye development and which may lead to treatment for relatede eye disorders or diseases. Sequences AAA46520-21
                                                                                                                                                                                                                                                                                         New isolated mammalian retinal calcium channel gene, used to develop products for the diagnosis and treatment of incomplete congenital stationary night blindness and related disorders.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human biallelic marker upstream amplification primer SEQ ID NO:4109.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human genome; biallelic marker; high density disequilibrium map; genomic map; haplotype; phenotype; polymorphic base; genotyping; haplotyping; hybridisation; identification; characterisation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     amplification; single nucleotide polymorphism; SNP; PCR primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0.8%; Score 13.8; DB 1; Length 20; 88.2%; Pred. No. 1e+03; Live 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Seguence 20 BP; 1 A; 6 C; 2 G; 11 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                 Disclosure; Page 15; 55pp; English
                                                                                                                                                         (UYTE-) UNIV TECHNOLOGIES INT INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1698 TTACTCTCTGCCTACCT 1714
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        BP
                                                                     99WO-CA000514
                                                                                                               98US-0087635P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAZ69753 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first entry)
                                                                                                                                                                                                       Bech-Hansen T, Naylor MJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  15; Conservative
                                                                                                                                                                                                                                                 WPI; 2000-097327/08.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              diagnosis; ss.
                                                                     02-JUN-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      21-APR-1998;
23-NOV-1998;
                                                                                                               02-JUN-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  409954500-A2
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                         09-DEC-1999,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           10-SEP-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAZ69753;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RESULT 1489
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAZ69753/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Matches
      ò
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ;
0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAX91991-X97517 represent PCR primers used to amplify open reading frames and other nucleic acid sequences from the genome of Chlamydia pneumoniae (see AAX91990). C. pneumoniae causes respiratory disease such as pneumonia and bronchitis and is thought to be a contributing factor in heart disease, sarcoidosis, sinusitis, purulent citis media, erythema nodosum or pharyngitis. The polypeptides encoded by the open reading frames of the C. pneumoniae genome (see AAY3454- AAY35879) can be used nucleotides sequences can also be used as immunogenic compositions, especially where the vector directs the expression of a neutralising epitope of C. pneumoniae
                                                                                                                      Respiratory disease; pneumonia; bronchitis; heart disease; sarcoidosis; sinusitis; purulent otitis media; erythema nodosum; pharyngitis; vaccine; neutralising epitope; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Retinal calcium channel, RCC gene, alphalF-subunit, retinal disorder; myopia, nystagmus; strabismus; calcium-regulated development pathway; eye disorder; human; EST; expressed sequence tag; CSNB; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0.8%; Score 13.8; DB 1; Length 20;
88.2%; Pred. No. 1e+03;
ative 0; Mismatches 2; Indels
                                                                                   PCR primer used to amplify an ORF of Chlamydia pneumoniae.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 20 BP; 6 A; 6 C; 6 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human EST JRL4Al amplifying forward primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Genome sequence of Chlamydia pneumoniae.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Page 1480; Disclosure; 1912pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1468 CTGGGGGGGGGTCCA 1484
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAZ46520 standard; DNA; 20 BP
                                                                                                                                                                                                                                                                                                                                                                                                                           97FR-00014673.
98US-0107078P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CTGCGAGAGCGGATCCA
                                            (first entry)
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                                                                                                                                                                                                                          Synthetic.
Chlamydophila pneumoniae.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 1999-357842/30.
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Matches 15; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (GEST ) GENSET
                                                                                                                                                                                                                                                                                           WO9927105-A2
                                                                                                                                                                                                                                                                                                                                                                                 20-NOV-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                  04-NOV-1998;
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                                            13-SEP-1999
                                                                                                                                                                                                                                                                                                                                     03-JUN-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Griffais R;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAZ46520;
AAX92036
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RESULT 1488

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Example 1; Page 30; 94pp; Japanese.

(GEST) GENSET

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AAZ65654 to AAZ69578 represent human biallelic markers from the present invention, which contain a polymorphic base at position 24 of their nucleotide sequences. AAZ69579 to AAZ7440 represent amplification primers for the biallelic markers. The biallelic markers of the invention have a variety of uses: they can be used for high density mapping of the human genome, and in complex association studies and haplotyping studies which are useful in determining the genetic basis for disease states. Compositions and methods of the invention can also be useful for the identification of the targets for the development of pharmaceutical agents and diagnostic methods, as well as the characterisation of the differential efficacious responses to and side effects from pharmaceutical agents acting on a disease as well as other treatment.

N. B. The SEQ ID NOS 2852, 2913, 2974, 3035, 3096, 3157, 327, 3297 and 3367, are not actually given a sequence in the Sequence Listing from the
                                                                                                         Novel biallelic markers used to construct a high density disequilibrium
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Serine protease BSSP6, useful in detecting homologs, mutants and polymorphic variants as markers for diagnosis of Alzheimer's disease, epilepsy, cancer, inflammation, infertility and prostate hypertrophy, using blood or other tissues.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 BSSP6; serine protease; human; hBSSP6; mouse; mBSSP6; brain; diagnostic marker; antibody; transgenic animal; Alzheimer's disease; epilepsy; cancer; inflammation; infertility; pancreatitis; prostatic hypertrophy; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human serine protease BSSP6 (hBSSP6), RACE PCR primer, SEQ ID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 13.8; DB 1; Length 20;
Pred. No. 1e+03;
0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Kominami K, Yamaguchi N, Mitsui S;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 20 BP; 2 A; 3 C; 6 G; 9 T; 0 U; 0 Other;
                                  Chumakov I;
                                                                                                                                                                Claim 8; Page 1107; 2745pp; English,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1076
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAA61782 standard; DNA; 20 BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1060 ATCCCAACAAGACATA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Matches 15; Conservative
                                  Blumenfeld M,
                                                                                                                            map of the human genome
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2000-400067/34.
                                                                      WPI; 2000-013267/01
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Uemura H, Okui A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                            present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WO200031257-Al.
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                                  Cohen D,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAA61782;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RESULT 1490
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The invention relates to novel serine proteases designated BSSP6

(AAB11712-B11714), and to nucleic acids encoding them (AAA61763-A61765).

The invention also relates to vectors and transformants comprising BSSP6

nucleic acids; transgenic animals in which the expression level of BSSP6

can be varied; and an mBSSP6 knockout mouse. The invention additionally

can be varied; and an mBSSP6 Anothodies and methods of production of such

antibodies, methods of BSSP6 detection using the antibodies, and the use

of BSSP6 proteins or fragments as diagnostic markers for certain medical

conditions. Nucleotides encoding BSSP6 were initially isolated in a human

brain cDNA library using degenerate PCR primers (AAA6179-A61796) based

on conserved regions of serine proteases. The BSSP6 serine proteases and

nucleotides encoding them are useful in detecting homologues, mutants and

conditions such as Alzheimer's disease. epilepsy, cancer, inflammation,

infertility and prostatic hypertrophy. Sequences AAA61768 and

crepresent PCR primers used in the exemplifications of the invention.

Primers AAA61775-A61784 and AAA61763, AAA61765, while primers AAA61785-

AA1792 were used to isolate and amplify murine BSSP6 cDNA (AAA61764).

Primers AAA61778 were used to construct plasmids used in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    14-3-3 sigma, HME1; stratifin; p53; diagnosis; cancer; psoriasis; polyp; psoriasis; wart; inflammatory disease; proliferation; ss; PCR primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Use of 14-3-3 sigma polypeptides and nucleic acids for the diagnosis or treatment of cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PCR primers AAX89470-X89471 are used to screen a BAC library for the
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0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 13.8; DB 1; Length 20;
Pred. No. 1e+03;
0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PCR primer used to screen a BAC library for 14-3-3 sigma.
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 20 BP; 2 A; 7 C; 3 G; 8 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Kinzler KW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example 3; Page 33; 73pp; English
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0.8%;
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98US-00210748
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         44 GAGGACCAGCAGTGTGA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        20 GAGCACCAGAAGTGTGA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DNA; 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        15; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAX89471 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   L5-FEB-2000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              18-DEC-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                     invention
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10017621-381.rng

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presence of a 14-3-3 sigma nucleotide sequence. 14-3-3 sigma is a member of the 14-3-3 protein family and is also known as HME1 or stratifin. 14-3-3 sigma expression is regulated by p53 and exogenous expression of 14-3-3 sigma results in G2 block. The 14-3-3 sigma nucleotide and amino acid sequences are used in the invention to develop agents for the diagnosis, susceptibility determination and treatment of cancer. The amino acid captures can be used in method for suppressing the growth of tumour cells. The 14-3-3 sigma polypeptics can mediate cell cycle arrest upon damage to cellular DNA. 14-3-3 sigma probes can be used for diagnosing, testing susceptibility to or treating cancers and identifying agents for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Antisense compound specifically hybridizing and inhibiting the expression of human Jun N-terminal kinase kinase-2 is useful for treating infection,
                                                                                                                                                         treating cancers. They can also be used to treat other proliferative diseases, e.g. psoriasis, polyps, warts, and inflammatory diseases. The 14-3-3 sigma antisense oligonucleotides can be used for promoting the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human; jun N-terminal kinase kinase-2; JKK-2; modulation; tumour;
antiinflammatory; cytostatic; antiinfectious; infection; inflammation;
detection; antisense therapy; phosphorothiaote; ss.
                                                                                                                                                                                                                                                                                                        Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human jun N-terminal kinase kinase-2 antisense oligonucleotide #33
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                                                                                                                                                                                                                                                                       DB 1; Length 20;
                                                                                                                                                                                                                                                                                                      2; Indels
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                                                                                                                                                                                                                                          Sequence 20 BP; 3 A; 9 C; 3 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                     Score 13.8; DB 1
Pred. No. 1e+03;
0; Mismatches
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                                                                                                                                                                                                           proliferation and growth of cells
                                                                                                                                                                                                                                                                                                     0;
                                                                                                                                                                                                                                                                                                                                    843 TGAGTACCTGGACAAGG 859
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                                                                                                                                                                                                                                                                       0.8%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                              AAA29848 standard; DNA; 20
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/*tag=
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Monia BP, Cowsert LM;
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Best Local Similarity
Matches 15; Conserv
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modified_base
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               24-JUN-1999;
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The present invention describes an antisense compound (I) of 8-30 nucleobases, specifically hybridising to, and inhibiting expression of, human jun N-terminal kinase kinase-2 (VKK-2). Also described is a method of inhibiting the expression of human JKK-2 in human cells or tissues, comprising contacting the cells or tissues, with (I), in vitro. (I) has antiinflammatory, cytostatic and antiinfectious activities. (I) is useful

Claim 3; Col 40; 31pp; English.

inflammation and tumor.

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which are components of the nomega bydroxy, an equation and the components of the DNA sequences (AAA10566-A30577) respectively reopticalis ATCC 20366. The DNA sequences (AAA10566-A30577) respectively arodycohrome P450 NADPH oxidoreductases CPRA and CPRB (AAY90596, CYP52A2A, CYP52A3B, CYP5A3AB, CYP5AAB, CYP5AB, CYP5ABB, CYP5ABB
                                                                                                                                                                                                                                                                                                         0
for inhibiting the expression of JKK-2 in human cells or tissues and prevents or delays infection, inflammation or tumour formation associated with altered expression of JKK-2. (I) is also useful for detecting the levels of JKK-2 in a sample. The present sequence represents a phosphorothioate antisense oligonucleotide for human JKK-2, from the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   omega hydroxylase complex; omega-oxidation; fatty acid; alkane; alpha-omega-dicarboxylic acid production; duntitative competitive reverse transcription-PCR; QC-RT-PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Cytochrome P450 nicotine adenine dinucleotide phosphate oxidoreductase and cytochrome P450 monooxygenase nucleic acids and encoded proteins, useful for overproducing dicarboxylic acids.
                                                                                                                                                                                                                                                                                                      Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The invention relates to 12 novel genomic DNA sequences and proteins
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Cytochrome P450; NADPH reductase; monooxygenase; CYP52A; CPR; POX;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     C. tropicalis CYP52A5A/CYP52A5B QC-RT-PCR primer, SEQ ID NO:47.
                                                                                                                                                                                                                                                                                                         ;
0
                                                                                                                                                                                                                                            0.8%; Score 13.8; DB 1; Length 20; 38.2%; Pred. No. 1e+03; ve 0; Mismatches 2; Indels
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                                                                                                                                                                                               Sequence 20 BP; 2 A; 4 C; 10 G; 4 T; 0 U; 0 Other;
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Gleeson M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 11; Page 44; 200pp; English,
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                                                                                                                                                                                                                                                                                              0;
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                                                                                                                                                                                                                                                                                                                                                           974 ACCGAGACCTCAAGCCC 990
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Candida tropicalis; ATCC20366.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ВР
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                                                                                                                                                                                                                                                                           88.2%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAA30532 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
                                                                                                                                                                                                                                                                                                         Conservative
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Tang M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (revised)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2000-317711/27.
                                                                                                                                                                                                                                                                           Local Similarity
les 15; Conserv
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21-AUG-2000
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                                                                                                                                                                                                                                                  Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         RESULT 1493
                                                                                                                                                                                                                                                                             Best Loca
Matches
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     8888888
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fatty acids via a novel quantitative competitive reverse transcriptionCPC (CC-RT-PCR). This assay quantifies the amount of target mRNA in a
sample and may be used for discriminating members of a gene family, such
as the CYP gene family. Organisms containing the target gene are cultured
on an organic substrate which causes upregulation of that gene. The total
RNA is then extracted and mixed with a known amount of competitor RNA,
which is similar to the target mRNA but has fewer nucleotides. RT-PCR
teactions are performed using increasing amounts of competitor RNA and
the point at which the amount of the corresponding target DNA is equal to
the amount of the amount of the corresponding competitor DNA is determined. The CPR and
CVP nucleic acids may be transformed into a suitable host so that the
host overspresses the corresponding proteins. Such host cells will
coverproduce dicarboxylic acids. The dicarboxylic acids thus produced find
spplication as thermoplastics, plasticising agents, lubricants, hydraulic
fluids, agricultural chemicals, pharmaceuticals, dyes, surfactants,
cathesives and fragrances cale production of industrially useful
dicarboxylic acids. Sequences AAA10522-A30543 represent QC-RF-PCR primers
cused in an exemplification of the invention to amplify the Candida
cused in an exemplification of the invention to amplify the Candida
cused in an exemplification of the invention pox gene target mRNA. BP; 7 A; 3 C; 9 G; 1 T; 0 U; 0 Other; Updated on 15-SEP-2003 to standardise OS field) Sequence 20

0.8%; Score 13.8; DB 1; Length 20; 2; Pred. No. 1e+03; 0; Mismatches 1010 AGAGGGGAGAGCTCAAG 1026 2 AGAGGCCAGGGCTCAAG 18 88.28; Local Similarity 88.2 nes 15; Conservative Query Match Best Loca Matches 엄 à

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Gaps

; 0

Indels

AAA78243 standard; DNA; 20 BP. (first entry) 16-NOV-2000 AAA78243; RESULT 1494 AAAAAB243
IID AAAAA
XX AAAA
XX AAAA
XX AAAA
XX Inmun
XX Inmu
XX Inmu
XX AAII
X

Anti-human Fas antibody CH11 H chain cDNA specific primer SHR-16. Antirheumatic agent; immunoglobulin M; IgM; apoptosis inducer; immunosuppression; autoimmune disease; treatment; rheumatism; anti-Fas antibody; primer; ss.

Synthetic

JP2000154149-A. 06-JUN-2000 98JP-00264598. (SANY) SANKYO CO LTD. 18-SEP-1998;

99JP-00263984.

17-SEP-1999;

WPI; 2000-454476/40.

Anti-human Fas humanizing antibody-containing antirheumatic agents

Disclosure; Page 16; 109pp; Japanese.

The present invention relates to antirheumatic agents which comprise as active ingredients an immunoglobulin M (1gM) protein. The 1gM protein does not include a J segment, has apoptosis inducing activity, and consists of a light and heavy chain polypeptide produced synthetically. The agents of the invention exhibit antirheumatic and immunosuppressive activity and can be used to treat autoimmune diseases, especially rheumatism. The 1gM molecule used in the invention has human Fas-antigen binding properties. Included in the invention are nucleotide sequences of

ö the IgM light and heavy chains (see AAA78267-A78272) and the corresponding protein sequences (see AAB12913-B12918 and AAB12919), and nucleotide sequences of the humanised anti-human Fas Ig CH11 (see AAA78202-A78206) and protein sequences (see AAB12908-B12910). Also included are anti-human Fas antibody CDR peptides (AAB12902-B12907). Primers specific for the anti-human Fas antibody, light, heavy and kappa chains used in the invention are represented by sequences AAA78213-A7826. Primers used for sequencing the human Ig DNA used in the invention are represented by sequences AAA78213-A78337, while humanised anti-Fas Ig DNA sequencing primers are represented by sequences AAA78212-A78338 and AAA78337. Primer represented by sequences PAA78321-A78338 and AAA78338-A78367. Primer represented by sequences PAA78321-A78334 and AAA78338-A78367. Primer represented by sequences PAA78321-A78334 and AAA78338-A78367. Primer represented by sequences PAA78321-A78334 and AAA78338-A78367. Primer broduction of the agent of the invention Gaps ; 0 0.8%; Score 13.8; DB 1; Length 20; Indels Sequence 20 BP; 2 A; 8 C; 2 G; 8 T; 0 U; 0 Other; 5; Pred. No. 1e+03;); Mismatches ., 1452 TCCATTCTTCCTCAGTC 1468 20 88.2%; 4 TCCATTCCTCCTCTGTC 15; Conservative Best Local Similarity Matches 15; Conserv Query Match g à

BP. AAZ59944 standard; DNA; 20 (first entry) 08-MAY-2000 AAZ59944; RESULT 1495 AAZ59944/c

expressed sequence tag; Drug exposure; drug abuse; gene expression; EST; expressed sequidentification; tolerance; addiction; therapy; screening; cellular response; ethanol; expression analysis; Northern blot; dopamine beta-hydroxylase; DBH; norepinephrine; reverse transcriptase-PCR; RT-PCR primer; ss. Human dopamine beta-hydroxylase (DBH) PCR primer, SEQ ID NO:2.

Homo sapiens.

99WO-US013839. 98US-0090268P. 22-JUN-1999; 22-JUN-1998; 21-JUN-1999; 29-DEC-1999.

WO9967267-A1

Lai C, Lockhart DJ; CALIFORNIA (REGC) UNIV Miles MF,

WPI; 2000-147195/13

Novel methods for evaluating an organisms response to alcohol used evaluate drug treatment and identifying susceptibility to alcohol.

Example 3; Page 68-69; 98pp; English.

t C

The invention relates the identification of genes whose expression levels are altered by chronic exposure to one or more drugs of abuse (e.g., ethanol, stimulants, opiates). The methods of the invention monitor the response of a cell to a drug of abuse, and comprise contacting the cell with the drug of abuse, and detecting the expression of one or more of 218 expressed sequence tags (ESTs) via the use of probes that specifically hybridise to the ESTs. The methods are used to identify genes whose expression levels are altered by chronic or acute exposure to one or more drugs of abuse. The identification of genes whose regulation is altered in alcohol tolerance and/or addiction provides a valuable tool

of a pituitary trophic hormone gene promoter; (4) identify a human afflicted with a disease, disorder, or condition caused by altered expression of Lhx3 or altered level of binding of Lhx3 to a nucleic acid; and (5) detect a mutation in a Lhx3 allele in a human. The coding region of human Lhx3 has been genomically mapped to the subtelomeric region of human Lhx3 is also known as P-LIM or LIM-3. The present sequence represents a PCR primer used in the amplification of human Lhx3, which is used in an example from the present invention

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the response to one or more drugs of abuse. Evaluation of the
            nature of this response provides information useful in designing the another or patient to drugs in a medical context. Monitoring the expression or patient to drugs in a medical context. Monitoring the expression of identified genes and/or ESTS provides a mechanism by which test agents can be screened for the ability to alter or modulate the response of the organism to drugs of abuse. Sequences AASS9944-259951 represent reverse transcriptase-PCR (RT-PCR) primers used to amplify 4 CDAM hybridisation probes from SH-SYSY-AH1861 human neuroblastoma cell expression in control and ethanol-treated SH-SYSY-AH1861 cells in an exemplification of the present invention. The genes whose expression was analysed were dopamine beta-hydroxylase (DBH) and sodium dependent
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The present invention describes an isolated nucleic acid (I) encoding a mammalian Lhx3. (I) is used in assays to: (1) detect and quantify the presence and level of expression of Lhx3, Lhx3a or Lhx3b, in a sample; (2) identify a compound that affects expression, the level of expression, or the activity of Lhx3, Lhx3a, or Lhx3b in a cell; (3) identify a compound that affects binding or Lhx3 to nucleic acid or Lhx3 induction
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     a l
                                                                                                                                                                                                                                                      norepinephrine transporter (NET), both of which are involved in corepinephrine metabolism; delta-like protein (DEK); and monocyte chemoattractant pepride 1 (MCP-1). These genes are thought to be important targets of ethanol. Primers AAZ59944-Z59945 were used to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New isolated nucleic acid encoding mammalian Lhx3 for identifying with a disease, disorder, or condition caused by an altered level expression or binding of Lhx3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Lhx3; LIM-3; P-LIM; identification; characterisation; diagnosis; chromosome 9; pituitary disease; subtelomeric region; mutation; pituitary trophic hormone gene promoter; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                             ó
                                                                                                                                                                                                                                                                                                                                                                                                                  0.8%; Score 13.8; DB 1; Length 20; BB.2%; Pred. No. 1e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Rhodes SJ, Bridwell JL, Meier BC, Parker GE, Price JR;
Showalter AD, Sloop KW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                         2; Indels
                                                                                                                                                                                                                                                                                                                                    generate the dopamine beta-hydroxylase (DBH) probe
                                                                                                                                                                                                                                                                                                                                                                           Sequence 20 BP; 3 A; 8 C; 5 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                    Pred. No. 1e+03;
); Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human Lhx3 exon 6 PCR primer SEQ ID NO:113.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (ADRE-) ADVANCED RES & TECHNOLOGY INST.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 6; Page 169; 239pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                         ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                400 GIGCAGICTCCAGIGAG 416
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            22-FEB-2000; 2000WO-US004424.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Grecagrasccagreas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAA92148 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2000-594085/56.
                                                                                                                                                                                                                                                                                                                                                                                                                                  Best Local Similarity
Matches 15; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WO200050868-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   22-FEB-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 04-JAN-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       31-AUG-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAA92148;
                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1496
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The present invention describes a radiation hybrid map of the dog (Canine familiaris) genome comprising the genome location of a marker selected from AAA66139 to AAA66942. The radiation hybrid map is useful for identifying and localising dog genes, since it covers approximately 80 % of the dog genome and provides a dense map integrating different types (i.e. Type I and Type II) of markers. The map and the dog genome markers (or complementary sequences) are especially useful to identify genes responsible for phenotypic and behavioural traits in dogs, to identify morbid genes, to analyse diseases and identify implicated genes in such diseases and thair alleles, and to study dog pedigrees. They may also be useful for isolating corresponding human gene sequences e.g. genes
                                                                                                                                                                                                                      Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Dog; genome; genomic marker; radiation hybrid map; identification; chromosome location; gene marker; polymorphic microsatellite marker; phenotype; behaviour; pedigree; ss.
                                                                                                                                                                                                                    .
0
                                                                                                                                                                                   DB 1; Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 13.8; DB 1; Length 20; Pred. No. 1e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Jog genomic marker oligonucleotide sequence SEQ ID NO:746.
                                                                                                                                                                                                                    2; Indels
                                                                                                                                                  Sequence 20 BP; 2 A; 4 C; 9 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 20 BP; 6 A; 4 C; 7 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                 0.8%; Score 13.8; DB 1
18.2%; Pred. No. 1e+03;
ive 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 1; Page 85; 87pp; English.
                                                                                                                                                                                                                                                    389 CCTCGGATGAGGTGCAG 405
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                                                                                                                                                                                                                                                                                                                                                                       BP
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                                                                                                                                                                                 Query Match 0.8%;
Best Local Similarity 88.2%;
Matches 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     involved in genetic diseases
                                                                                                                                                                                                                                                                                                                                                                     AAA66884 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                      (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Andre C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2000-387821/33.
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Canis familiaris.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WO200029615-A2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           25-MAY-2000.
                                                                                                                                                                                                                                                                                                                                                                                                      AAA66884;
                                                                                                                                                                                                                                                                                                                                       RESULT 1497
                                                                                                                                                                                                                                                                                                                                                     AAA66884
ID AAA
8888888888888
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Matches

ð ď RESULT 1498

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cubstances (1) containing recombinantly prepared subunits (SU) based on amino acids (aa) comprising: (i) at least two modified SU with one difference; and/or (ii) one or more modified SU with at least two differences; and/or (ii) one or more modified SU with at least two differences; and (iii) (optionally) modified SU. The various SU are combined in a mosaic fashion to form (A) in which (I) can be enapsulated. (A) Are used to deliver (I) specifically to cells, by human immune deficiency virus, tumors and a wide range of inherited of seases such as hemophilia, muscular dystrophy or cystic fibrosis. Composition. SU can be motified to impart a wide variety of selected composition. SU can be modified to impart a wide variety of selected composition. SU can be modified to impart a wide variety of selected composition. SU can be modified to impart a wide variety of selected communogenicity. (A) do not require extensive testing to ensure that they are safe (contrast replication-deficient viruses), also SU can be prepared in very pure form and are easily labeled fluorescently (for quality control or localization). This sequence represents a PCR primer cused in the production of a Listerial monocytogenes listeriolysin LLO variant which is used to illustrate the method of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Macrophage migration inhibitory factor; MMIF; antisense; neurological; hyperproliferation; nootropic; antihormonal; immunosuppresive; human; antiinflammatory; cytostatic; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Transport system for compounds, useful e.g. in gene therapy, comprises mosaic-like assembly of different protein subunits able to encapsulate
                                       Transport system; gene therapy; infection; tumor; ss; LLO; PCR primer; human immune deficiency virus; hemophilia; muscular dystrophy; capsid; cystic fibrosis; virus-like particle; cell targeting; listeriolysin O.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             This invention describes a novel transport system (A) for molecular substances (I) containing recombinantly prepared subunits (SU) base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0.8%; Score 13.8; DB 1; Length 20; 38.2%; Pred. No. 1e+03;
L. monocytogenes listeriolysin O variant LLO PCR primer #3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human MMIF mRNA inhibiting antisense oligo ISIS #112711.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 20 BP; 3 A; 13 C; 3 G; 1 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                     Esser
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                                                                                                                                                                                                                                                                                                                                                                                     Schmidt
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Н
                                                                                                                                                                                                                                                          03-NOV-2000; 2000WO-EP010876.
                                                                                                                                                                                                                                                                                                    99DE-01052957
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ilarity 88.2%;
Conservative
                                                                                                                                                                                                                                                                                                                                           (ACGT-) ACGT PROGENOMICS AG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           230 GTGGTGGTGGTGGCGGC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         standard; DNA;
                                                                                                                                 Listeria monocytogenes
                                                                                                                                                                                                                                                                                                                                                                                   Rudolph R,
                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2001-316433/33
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                                                                                                                                                                      WO200132851-A2
                                                                                                                                                                                                                                                                                                    03-NOV-1999;
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                                                                                                                                                                                                                  10-MAY-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              compounds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAH23201;
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Matches
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      ó
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                830 Primers useful for synthesizing full length cDNA clones and their use
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The invention relates to primers for synthesising full length cDNA clones. 830 cDNA molecules encoding a human protein have been isolated and nucleotide sequences of 5'- and 3'-ends of the cDNA molecules have been determined. Primers for synthesising the full length cDNA are useful for clarifying the function of the protein encoded by the cDNA. The full length clones were obtained by construction of full length enriched cDNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          libraries that were synthesised by the oligo-capping method. The primers enable the production of the full length cDNA easily without any special methods. The present sequence is a primer work of the full length cDNA easily without any special
                                                                                                                                                                                                                                                                                                                                                                   Human; full length cDNA; cDNA synthesis; oligo-capping; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       methods. The present sequence is a primer used to amplify a human cDNA clone provided in the invention
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        Gaps
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T, Koga 1
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8.2%; Pred. No. 1e+03;
ve 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example 18; Page 132; 1380pp + Sequence Listing; English.
        Indels
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K, Kojima S, Otsuki
                                                                                                                                                                                                                                                                                                                            Human cDNA clone-specific primer, SEQ ID NO: 4416.
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        2
        Mismatches
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a T, Nagai
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11-JAN-2000; 2000JP-00118774.
02-MAY-2000; 2000JP-00183765.
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                                                                                          4 GGAAGAGGTGCAATCTC
                                                                                                                                                                                                 AAK95171 standard; DNA; 20
                                                  393 GGATGAGGTGCAGTCTC
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Best Local Similarity 88.4.
Best Local Similarity 88.4.
                                                                                                                                                                                                                                                                                    (first entry)
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        15; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      in genetic manipulation.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Wakamatsu A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                     EP1130094-A2
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The invention relates to antisense oligonucleotides 8-30 nucleotides in length targeted to a nucleic acid molecule encoding macrophage migration inhibitory factor (MMF), where the antisense compound specifically hybridizes with and inhibits the expression of MMIF. The antisense nucleotides are useful for the treatment of a disease or condition associated with MMIF such as neurological, hormonal, immune, inflammatory or hyperproliferative disorder. Sequences AAH21191-268 represent chimeric antisense phosphorothioate oligonucleotides used for inhibition of human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Vaccine; cytostatic; virucidal; bactericidal; fungicidal; anti-parasitic; immunostimulatory; tumour; viral infection; bacterial infection; fungal infection; parasitic infection asthma; infection infection; manne deficiency; phosphorothicate; ss.
                                                                                                                                                                                                                                                                    New antisense compound(s) are useful to inhibit a nucleic acid molecule encoding macrophage migration inhibitory factor.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0.8%; Score 13.8; DB 1; Length 20;
88.2%; Pred. No. 1e+03;
tive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 20 BP; 7 A; 4 C; 8 G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Immunostimulatory nucleic acid #929.
                                                                                                                                                                                                                                                                                                               Claim 3; Page 82; 105pp; English
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       25-SEP-1999; 99US-0156113P.
27-SEP-1999; 99US-0156135P.
23-AUG-2000; 2000US-0227436P.
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                                                                                                                     16-JAN-2001; 2001WO-US001475.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (IOWA ) UNIV IOWA RES FOUND. (COLE-) COLEY PHARM GMBH.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   39 GGCAGGAGGACCAGCAG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GGCAGAGGACCAGGAG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAF99813 standard; DNA; 20
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Best Local Similarity 88.2
Matches 15; Conservative
                                                                                                                                                                                                          Cowsert LM,
                                                                                                                                                                             (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                 antisense phosphorotl
MMIF mRNA expression
                                                                                                                                                                                                                                      WPI; 2001-451899/48.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WO200122972-A2.
                                                          WO200153317-A1
                                Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   12-JUN-2001
                                                                                        26-JUL-2001
                                                                                                                                                                                                          Murray SF,
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                 Synthetic
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The present invention relates to a method for stimulating an immune response. The method comprises administering an immunostimulatory nucleic acid to a non-rodent subject in sufficient quantity to stimulate an immune response. The present sequence is one such immunostimulatory
                                                                                                                                                                                                                                      nucleic acid. The immunostimulatory nucleic acids can be pyrimidine rich (Pyritch) or thymidine (T) rich. The method is used to vaccinate subjects against tumour antiqens, viral antiqens (e.g. texperviridae, retroviridae and/or orthomyxoviridae), bacterial antiqens (e.g. texpolasma, haemophilus, campylobacter, clostridium, Escherichia coli and/or staphylococcus), fungal antiqens and/or parasitic antiqens. The method is also useful for preventing cancer, asthma, infectious disease, allergy or fimume deficiency. The present sequence can also be used to redirect a Th2 to a Th1 immune response and to activate immune cells. Note: the present sequence may have a phosphorothioate backbone
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Fascin; regulatory sequence; human; dendritic cell; antiviral; tumor; antibacterial; antifungal; antiparasitic; anti-allergic; neurological; immunomodulatory; apoptotic; expression regulator; vaccine; allergen; Creutzfeld-Jakob disease; Alzheiner's disease; gene therapy; autoimmune disease; transplant rejection; primer; ss.
                                                                                 Vaccinating against tumors, infectious diseases, allergies and asthma using immunostimulatory Py-rich and TG nucleic acids.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New regulatory sequences from the fascin gene, useful for providing
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0.8%; Score 13.8; DB 1; Length 20; 88.2%; Pred. No. 18+03; ive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 20 BP; 2 A; 5 C; 6 G; 7 T; 0 U; 0 Other;
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                                                                                                                                      Claim 101; Page 58; 338pp; English
              Vollmer J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1547 GCCTTCGGTCTTCGTCG 1563
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ВР
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 GCCTTCGATCTTCGTTG 17
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02-MAR-2000; 2000DE-01010188.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAH48588 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match 0.8
Best Local Similarity 88.2
Matches 15; Conservative
              Schetter C,
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                                                 WPI; 2001-273485/28
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ROSS R.
BROS M.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Homo sapiens
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            Krieg AM,
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(ROSS/)
(ROSS/)
(ROSS/)
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Gaps

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dendritic cell-specific expression of e.g. antigens, e.g. for vaccination
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                                                                                                                                                                                                                                 allergic, neurological, immunomodulatory and apoptotic activity. (A) are used to regulate expression of antigens, immunoregulators, antisense sequences etc. in DC-specific fashion. Recombinant DNA, vectors and host cells that contain (A) are useful: (i) in vaccines against viruses, bacteria, fungi, parasites, umors, allergens and plaques in Creutzfeld-Jakob and Alzheimer's disease; and (ii) for gene therapy of tumors,
                                                                                                                                                                                                                                                                                                                                                                                                                    allergies, infections, autoimmune diseases and transplant rejection. They
                                                                                                                                                                                                                                                                                                                                                                                                                                          can also be provide specific expression of antigens and immunoregulators in DC; for isolation and identification of cell factors and cis-elements from regulatory sequences that mediate DC-specific expression, to determine the degree of maturity of DC and to block transcription factors, by providing binding sites in DC. (A) provide DC-specific expression of nucleic acid under their control, allowing a more specific regulation of the immune response and eliminating the long and laborious purification of DC (since a complete leucocyte population may be transformed), including transformation in vitro. This sequence represents
                                                                                                                                                                         human fascin that provide specific expression in dendritic cells (DC) which have antiviral, antibacterial, antifungal, antiparasitic, anti-
                                                                                                                                                 (A)
                                                                                                                                                    sednences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 20 BP; 2 A; 4 C; 9 G; 5 T; 0 U; 0 Other;
                                                                                                                                                    novel regulatory
                                                                                         Claim 1d; Page 105; 117pp; German.
                              against tumors and infections.
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0.8%; Score 13.8; DB 1; Length 20;
88.2%; Pred. No. 1e+03;
ive 0; Mismatches 2; Indels
                                                                        986 AGCCCCAGAACCTGCTC 1002
                                                                                                        17 AGCCCCAGAACCCGCAC 1
Query Match 0.8
Best Local Similarity 88.2
Matches 15; Conservative
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Gaps

Canine retroviral PCR primer MLVIN5700+. AAC89128 standard; DNA; 20 BP. (first entry) Unidentified. 07-MAR-2001 AAC89128; RESULT 1503 AAC89128, E STAN SERVICE SERVICE

PCR primer; immunosuppressive; cytostatic; gene therapy; retrovirus; canine; autoimmune disease; haematopoietic malignancy; malignant tumour;

WO200070024-A2

23-NOV-2000

17-MAY-2000; 2000WO-EP004467

99EP-00401192 99EP-00401199 17-MAY-1999; 18-MAY-1999;

(FRSA-) ETAB FR DU SANG.

Darlix J; Corbine A, Ghernati I, Rigal D,

WPI; 2001-016224/02.

New infectious retrovirus isolated from a canine cell line, useful for producing medicaments to treat autoimmune diseases, hematopoietic malignancies or malignant tumors and in diagnosis and gene therapy.

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Claim 31; Fig 11; 131pp; English
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ô The present invention relates to a retrovirus of type C morphology, which sediments in a sucrose gradient at a density of 1.16-1.18 g/l. The retrovirus is infectious for canine cells and belongs to the oncovirinae group. The present sequence is a PCR primer for the retrovirus of the present invention. The retrovirus can be included in pharmaceutical compositions or medicaments to treat autonimune diseases, haematopoietic malignancies or malignant tumours, especially in humans. The retrovirus can also be used in gene therapy to introduce a transgene into an animal, The invention provides an agent comprising a pyrone compound or dihydroxy compound of specified formulae given in the specification. The agent is used for correcting gene expression in the specification. The agent is following genes may be corrected: IL-6, IL-10, hemeoxygenase-1, prostaglandin G/H synthase-2, macrophage inflammatory protein-1-alpha, IL-lbeta, TNF alpha, IL-1 receptor, macrophage inflammatory protein-1-alpha, macrophage inflammatory protein-1-beta, liver and activation-regulated chemokine, macrophage inflammatory protein-2-beta, marrix metalloproteinase-9, migration inhibitory factor-related protein-8, lyzozyme, GABA(A) receptor-associated protein, interferon-induced 17-Pyrone, gene therapy; antiinflammatory; gene expression; interleukin; hemeoxygenase-1; prostaglandin G/H synthase-2; RANTES; TNF alpha; p78; marcrophage inflammatory protein; chemokine; growth regulated protein-1; matrix metalloproteinase-9; migration inhibitory factor-related protein; lyzozyme; GABA(A) receptor-associated protein; interferon; SCO homolog-2; transketolase, adenosine A2a receptor, CD37 antigen properdin P factor; G-protein, Nef-associated factor-1; signal peptidase; PCR primer; ss. for correcting gene expression regulation error comprises pyrone Gaps Human GABA(A) receptor-associated protein specific primer GABArp-R. ô DB 1; Length 20; Indels Kato Sequence 20 BP; 3 A; 12 C; 2 G; 3 T; 0 U; 0 Other; 2; Sagawa H, 1e+03; Score 13.8; DE Pred. No. 1e+03 0; Mismatches Nishimura K, Example 6; Page 77; 93pp; Japanese. 0; 242 compound or dihydroxy compound BP. 11-JAN-2001; 2001WO-JP000082. 13-JAN-2000; 2000JP-00004989. 03-OCT-2000; 2000JP-00303711. 0.8%; 88.28; (TAKI) TAKARA SHUZO CO LID. AAH76258 standard; DNA; 20 226 GAGAGTGGTGGTGG 20 GAGAGCGGTGGGGGTGG (first entry) Best Local Similarity 88.2 Matches 15; Conservative Yamashita S, WPI; 2001-514436/56. especially a human WO200151480-A1 Homo sapiens. 29-0CT-2001 19-JUL-2001. AAH76258; Ľ Query Match RESULT 1504 Enoki Agent *55555555555 à P

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kDa/15-kDa protein, interferon-inducible protein p78, SCO homolog-2, transketolase, adenosine A2a receptor, CD37 antigen properdin P factor, regulator of G-protein signaling-2, Nef-associated factor-1, myeloid leukemia cell differentiation protein-1, signal peptidase complex, and also side-effects caused by them such as inflammation. Sequences AAH76220-76280 represent PCR primers used in the course of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PCR primers AAFB0165-66 were used to amplify the left-hand GAL7 promoter. The amplified sequence was used to express fusion proteins comprising a heavy chain variable region of an antibody isolated from llama, which was immunised with Malassezia furfur. M. furfur has been implicated in dandruff formation. The heavy chain variable region is conjugated to an active agent, and used to produce a composition for topical application, as shampoos and conditioners, skin care lotions, shower gels, etc., is useful for targeting an active agent to a site at which M. furfur is present for the treatment and prevention of dandruff hurfur is
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Composition for use in targeting active agent, especially antimicrobial agent to scalp for treating, preventing dandruff, has active agent conjugated to antibody capable of binding specifically to Malassezia
                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
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hair care; GAL7 promoter; PCR primer; ss.
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                                                                                                                                                                                                                                                                                                                                   0.8%; Score 13.8; DB 1; Length 20;
88.2%; Pred. No. 1e+03;
Live 0; Mismatches 2; Indels
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88.2%; Pred. No. 1e+03;
ative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                           Sequence 20 BP; 2 A; 4 C; 7 G; 7 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 917 TGTTCCTGTTCCAGCTG 933
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(UNIL ) UNILEVER NV.
(HIND-) HINDUSTAN LEVER LTD.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAF80165 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
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nes 15; Conserv
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Best Local Similarity Matches 15; Conserv

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New isolated polynucleotide useful for the identification of therapeutics in allergic diseases is new.
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88.2%; Pred. No. 1e+03;
tive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Denton RR, Duda A, Nandabalan K, Stephens JC;
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                                                                                                                                                                                                         Human IL4Ralpha gene PCR primer #48.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example 1; Page 61; 188pp; English.
                                                                                                                                                                                                                                                    allergic disease; PCR primer; ss
1472 GGGAGCGGATCCACAAA 1488
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              697
                             1 GGGAGGATCCAAAAA 17
                                                                                                            AAF69712 standard; DNA; 20 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                (GENA-) GENAISSANCE PHARM INC
                                                                                                                                                                                                                                                                                                                                                                               13-JUL-2000; 2000WO-US019094.
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                                                                                                                                                                        (first entry)
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                                                                                                                                                                                                                                     Polymorphism; human;
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                                                                                                                                                                                                                                                                                      Homo sapiens.
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                                                                                                                                                                         18-APR-2001
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                                                                                                                                          AAF69712;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ABZ72182;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Chew A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        RESULT 1507
ABZ72182/c
                                                                             RESULT 1506
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03-APR-2003 (first entry)

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The invention relates to isolated genes (Gene 216) from human chromosome 20p13-p12 and the proteins they encode. The nucleic acids and proteins may be used in the prevention, diagnosis and treatment of diseases associated with inappropriate Gene 216 expression. For example, the nucleic acids (or vectors) and proteins may be used to treat disorders associated with decreased expression by rectifying mutations or deletions in a patient's genome that affect the activity of gene 216 by expressing in a patient's genome that affect the activity of gene 216 by expressing in a control or complement or to supplement the patients own production of Gene 216 proteins. Additionally, the nucleic acids may be used to produce the secreted Gene 216 protein, by inserting the mucleic acids into a host complementary sequences may also be used as DNA probes in diagnostic assays to detect and quantitate the presence of similar nucleic acid sequences in samples and therefore which patients may be in need of restorative therapy. The Gene 216 protein may also be used as antigens in the production of antibodies against Gene 216 and in assays to identify modulators of Gene 216 expression and activity. The anti-Gene 216 antibodies may also be used as diagnostic and activity. The anti-Gene 216 antibodies may also be used as diagnostic agents for detecting the presence of Gene 216 proteins in samples (e.g. by enzyme linked immunosorbant assay or EUISA). Disorders that may be
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Isolated genes (Gene 216) from human chromosome 20p13-p12 and the proteins they encode, useful for the prevention, diagnosis and treatment of asthma, obesity and inflammatory bowel disease.
              Human, Gene 216; chromosome 20p13-p12; antiasthmatic; anorectic; antiinflammatory; gastrointestinal; gene therapy; vaccine; asthma; obesity; inflammatory bowel disease; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 20 BP; 4 A; 4 C; 7 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example 10; Page 149; 520pp; English
                                                                                                                                                                                                                                                                                                                                     (GENO-) GENOME THERAPEUTICS CORP.
                                                                                                                                                                                                                                                                                        13-APR-2000; 2000US-00548797.
                                                                                                                                                                                                                                              13-APR-2001, 2001WO-US012245.
                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2001-639428/73.
                                                                                                                                                      WO200178894-A2.
                                                                                                                                                                                                 25-OCT-2001
                                                                                                                                                                                                                                                                                                                                                                                  Keith T;
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C)
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prevented, diagnosed and/or treated by the above methods include, for example asthma, obesity and inflammatory bowel disease. The present sequence is that of a gene 216 related primer used in examples of the invention. The primers are used in the physical mapping of the gene
                                                                                                                                                                                                                                                                            Gaps
                                                                                            (ABZ72067-ABZ72088), polymorphism identifiction using single strand
                                                                                                               conformational polymorphism (SSCP) analysis (ABZ7209I-ABZ72184), sequencing (ABZ72185-ABZ72268) and genotyping (ABZ72317-ABZ72362)
                                                                                                                                                                                                                                                                            0
                                                                                                                                                                                                                               0.8%; Score 13.8; DB 1; Length 20;
                                                                                                                                                                                                                                                                            Indels
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                                                                                                                                                                                                                                                                                                                       538 CCCATCTTTGACAAGCC 554
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                                                                                                                                                                                                                                                  88.2%;
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                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                                                                                                                               Local Similarity
es 15; Conserv
                                                                                                                                                                                                                                                                                                                                                                 19
                                                                                                                                                                                                                                 Query Match
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Matches
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ABZ72122

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ABZ72122

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Life invention telature to landered genes (vene 210) if the invention intelature to landered genes (vene 210) if the intelature of the proteins they encode. The nucleic acids and proteins case may be used in the prevention, diagnosis and treatment of diseases associated with inappropriate Gene 216 expression. For example, the nucleic acids (or vectors) and proteins may be used to treat disorders associated with decreased expression by rectifying mutations or deletions in a patient's genome that affect the activity of gene 216 by expressing in a patient's genome that affect the activity of gene 216 by expressing contactive proteins or to supplement the patients own production of Gene 216 proteins. Additionally, the nucleic acids may be used to produce the secreted Gene 216 protein, by inserting the nucleic acids into a host complementary sequences may also be used as DNA probes in diagnostic assays to detect and quantitate the presence of similar nucleic acids and complementary sequences may also be used as DNA probes in need of restorative therapy. The Gene 216 protein may also be used as antigens in the production of antibodies against Gene 216 and in assays to identify modulators of Gene 216 expression and activity. The anti-Gene 216 antibodies may also be used as diagnostic and activity. The anti-Gene 216 antibodies may also be used as diagnostic and activity. The anti-Gene 216 antibodies may also be used as diagnostic and activity. The anti-Gene 216 antibodies may also be used as diagnostic appropriate and presence of Gene 216 proteins in samples (e.g. by enzyme linke defining tassay or ELISA). Disorders that may be proposed and/or events above method in an and/or events.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Isolated genes (Gene 216) from human chromosome 20p13-p12 and the proteins they encode, useful for the prevention, diagnosis and treatment of asthma, obesity and inflammatory bowel disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The invention relates to isolated genes (Gene 216) from human chromosome
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   prevented, diagnosed and/or treated by the above methods include, for example asthma, obesity and inflammatory bowel disease. The present sequence is that of a Gene 216 related primer used in examples of the invention. The primers are used in the physical mapping of the gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (ABZ72067-ABZ72088), polymorphism identifiction using single strand conformational polymorphism (SSCP) analysis (ABZ72091-ABZ72184), sequencing (ABZ72185-ABZ72268) and genotyping (ABZ72317-ABZ72362)
                                                                                            Human; Gene 216; chromosome 20p13-p12; antiasthmatic; anorectic; antiinflammatory; gastrointestinal; gene therapy; vaccine; asthma; obesity; inflammatory bowel disease; primer; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 20 BP; 5 A; 7 C; 4 G; 4 T; 0 U; 0 Other;
                                              Gene 216 SSCP detection primer SEQ ID NO 94.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example 10; Page 149; 520pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                (GENO-) GENOME THERAPEUTICS CORP.
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                                                                                                                                                                                                                                                                                                                                                                                                                     13-APR-2000; 2000US-00548797.
                                                                                                                                                                                                                                                                                                                                                           13-APR-2001; 2001WO-US012245
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nes 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2001-639428/73
                                                                                                                                                                                                                                                          WO200178894-A2.
                                                                                                                                                                                                                                                                                                             25-OCT-2001,
                                                                                                                                                                                                         Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Keith T;
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ID ABS7
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The present invention relates to a screening method for compounds for their ability to modify the binding of G-protein coupled receptor (GPCR) protein ZAQI and related proteins (human ZAQI, rat ZAQI (KZAQI), rZAQZ, human and mouse ISE (mISE) receptor, and mouse GPR73) to receptor protein and ligand are contacted in the presence or absence of the rest compound. The compounds are useful in a drug composition for the treatment, and prevention of digestive and central nervous system (CNS) disorders, including bowel inflammation, diarrhoea, constipation, food absorption disorders, Alzheimer's disease, Parkinson's disease and schizophrenia. The present sequence represents a PCR primer used in the
                                                                                                          G-protein coupled receptor; GPCR; ZAQ; human; ZAQC; ZAQT; rat; ZAQ1; rZAQ1; rZAQ2; mouse; 15E receptor; m15E; GPR73; Bv8 protein; digestive disorder; central nervous system disorder; CNS; diarrhoea; bowel inflammation; constipation; food absorption disorder; nootropic; Alzheimer's disease; Parkinson's disease; schizophrenia; laxative; antiinflammatory; antidiarrhoeic; neuroleptic; neuroprotective; PCR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Screening of compounds modifying the binding of G-protein coupled receptor protein ZAQ and related proteins to their ligands for use in treatment and diagnosis of digestive disorders.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PCR primer PV3 used to amplify HPV in human cervical cancer cells.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0.8%; Score 13.8; DB 1; Length 20; 88.2%; Pred. No. 1e+03; ative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Terao Y,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 20 BP; 3 A; 7 C; 3 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Watanabe T,
                                                                              Rat GPCR ligand Bv8 cDNA PCR primer RBv8-WR2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example 5; Page 127; 197pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              examples of the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Takatsu Y,
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                                                                                                                                                                                                                                                                                                                                                     01-FEB-2002; 2002WO-JP000852.
                                                                                                                                                                                                                                                                                                                                                                                    02-FEB-2001; 2001JP-00026820
                                                                                                                                                                                                                                                                                                                                                                                                                    (TAKE ) TAKEDA CHEM IND LTD.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      194/c
AAH77194 standard; DNA; 20
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(first entry)
                                               (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                     Masuda Y,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2002-627537/67.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Local Similarity
                                                                                                                                                                                                                                                                                  WO200262944-A2.
                                               27-NOV-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   07-AUG-2003
24-JAN-2002
                                                                                                                                                                                                                                                                                                                  15-AUG-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                15;
                                                                                                                                                                                                                   primer; ss
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S
                                                                                                                                                                                                                                                 Rattus sp.
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              ABS71117;
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Shintani Y;

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The sequence represents PCR primer PV3. The primer was used in the invention to amplify DNA from cells of the stable human cervical cancer cell line SiHa. containing on integrated copy of human papilloma virus (HPV) type 16 per human genome. The invention relates to a novel thermal cycling device for regulating the temperature of a biological sample on a flat substrate. The invention also includes an apparatus comprising the flat substrate for use in the thermal cycling device. The invention is useful for medical diagnosis of diseases such as AIDS, also for amplification of nucleic acids in biological samples. The invention has the advantage that it enhances operatively as the heat resisting element is directly coupled to the microscopic slide, and reduces costs as the use of a heat sink is eliminated. (Updated on 07-AUG-2003 to correct OS
                                                                                                                                                                                                                                                                                                                   Microscopic slide temperature control apparatus for medical diagnosis comprises coupling resistive heating element between the connection pads
Human, cervical cancer; human papilloma virus; PCR primer; PV3; SiHa; HPV; Thermal cycling; AIDS; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Rice; lesion formation inhibition; heat stress; agriculture; Sp17; transgenic; plant; horticulture; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Rice lesion inhibitor protein Sp17 coding sequence PCR primer #9.
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0
                                                                                                                                                                                                                                                   Arik M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0.8%; Score 13.8; DB 1; Length 20; 88.2%; Pred. No. 1e+03; ve 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                   Cibuzar GT, Schiller P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Seguence 20 BP; 2 A; 4 C; 6 G; 8 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                      provided at opposing ends of slide
                                                                                                                                                                                                                                                                                                                                                                                       Example 3; Col 27; 25pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1308 CAAGACATACAACTACC 1324
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                                                                                                                                                   99US-00432012.
                                                                                                                                                                                   99US-00432012.
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                                                                                                                                                                                                                                                   3lumenfeld M, Bar-Cohen A,
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Matches 15; Conservative
                                                                                                                                                                                                                  (MINU ) UNIV MINNESOTA
                                                  Human papillomavirus
                                                                                                                                                                                                                                                                                  WPI; 2002-009526/01.
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                                                                                   JS6300124-B1
                                                                                                                                                   32-NOV-1999;
                                                                                                                                                                                   02-NOV-1999;
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The polypeptide and its associated DNA sequence can be produced by recombinant techniques and is useful for preventing obesity, diabetes or alcoholism associated with SACI expression. The sequences are useful in screening for drugs and sweeteners. Recombinant cell lines and transgenic embryos may be used in screening for and identifying agents that induce or repress function of SACI. Predisposition to diabetes, obesity or alcoholism can be ascertained by testing any fluid or tissue of a human (such as blood, pancreas or tongue) for sequence variations of the SACI predisposition to diabetes, obsesty and/or alcoholism and may provide a diagnostic mark. The polynucleotide can be detected in a biological sample by contacting the DNA with a probe to form a hybridisation complex which is then detected. The sequences represent cDNA encoding human and mouse SACI polypeptides and PCR primers specific for the SCAI genes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human; Talin, antimicrobial; antiinflammatory; cytostatic; inhibitor; antisense gene therapy; infection; inflammation; Talin inhibitor; tumour; antisense oligonucleotide; phosphorothioate; ss.
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with altered preference for carbohydrates, other sweeteners or ethanol.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human Talin antisense phosphorothioate oligonucleotide SEQ ID NO:77.
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/note= "2'-methoxyethyl (2'-MOE) nucleotides"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           'note = "phosphorothioate backbone"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0.8%; Score 13.8; DB 1; 188.2%; Pred. No. 1e+03; tive 0; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 20 BP; 6 A; 3 C; 8 G; 3 T; 0 U; 0 Other;
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ABN89264;
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                                                                                                                                                                                                                                                                                                                                                                                                     The present invention provides the protein and coding sequences of rice lesion formation inhibitor Sp17. The protein improves the heat stress of the plant, and can be used in the development of new breeds of plants for agriculture and horriculture. The present sequence is a PCR primer used to isolate the coding sequence of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Novel isolated polypeptide comprising variant form of mouse or human SACI polypeptide, and is associated with altered preference for carbohydrates or other sweeteners, useful for preventing obesity, diabetes, alcoholism.
                                                                                                                                                            ALICE-Originated gene, Sp17, that inhibits lesion formation and is applicable in improving heat stress of plants thus leading to prevention of lesion formation, for developing new breeds of plants for agriculture and horticulture.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human; mouse; SAC1; carbohydrate; sweetener; ethanol; alcoholism; ss;
obesity; diabetes; transgenic embryo; body tissue; body fluid; pancreas;
blood; tongue; PCR primer; anorectic; antidiabetic; gene therapy;
protein replacement therapy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The invention relates to an isolated polypeptide, comprising a variant form of mouse or human SAC1 polypeptide. The variant form is associated
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88.2%; Pred. No. 1e+03;
tive 0; Mismatches 2; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 20 BP; 1 A; 6 C; 8 G; 5 T; 0 U; 0 Other;
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Tordoff MG;
      (NAAG-) NAT INST AGROBIOLOGICAL SCI
                                                                                                                                                                                                                                                                                                                                              Example 6; Page 47; 53pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 14; Page 90; 239pp; English
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28-JUL-2000; 2000US-0221419P.
10-NOV-2000; 2000US-0247443P.
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                                                                  Yano M, Yamanouchi
                                                                                                                            WPI; 2002-372312/40.
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10017621-381.rng

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The present invention describes an antisense compound (I), 16 to 30 bases in length targeted to specific base regions of a nucleic acid encoding human Talin. Also described: (a) an antisense compound up to 30 bases in length which inhibits the expression of human Talin; (b) a composition (II) comprising (I) or (a); and (c) inhibiting the expression of human tissues in human cells or tissues comprising contacting the cells or tissues in vitro with (I) or (a). (I) has antimicrobial, antiinflammatory and cytostatic activities, and can be used in antisense gene therapy and as a Talin expression inhibitor. (I) can be used: to inhibit the expression of human Talin in human cells or tissues; to prevent or delay
                                                                                                                                                                                                                                                                                                                                                                                              infection, inflammation or tumour formation; and in diagnostics, therapeutics, prophylaxis, and in research reagents and kits. The present sequence represents a human Talin antisense chimeric phosphorothioate oligonucleotide, having 2'-methoxyethyl (2'-MOE) wings of 5 nucleotides at the 5' and 3' ends and a 10 nucleotide deoxy gap in the middle, which is used in an example from the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 20 BP; 5 A; 7 C; 4 G; 4 T; 0 U; 0 Other;
                               Claim 14; Col 42; 46pp; English.
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0.8%; Score 13.8; DB 1; Length 20; 88.2%; Pred. No. 1e+03; ive 0; Mismatches 2; Indels 0; 1571 ACTCAGGCAGGCCAGCT 1587 20 4 ACTCTGGCAGGCCATCT Conservative Best Local Similarity Matches 15; Conser Query Match ð qq

ABS78535 standard; DNA; 20 (first entry) 13-DEC-2002 ABS78535 RESULT 1514
ABS78535
IX
AC ABS78535
XX
AC ABS7853
XX
XX
AC ABS7853
XX
XX
COT ABS7853
XX
COT ABS7

Angiogenesis inhibitory oligonucleotide #1019.

Angiogenesis inhibitor; ss; angiogenesis; solid tumour growth; tumour metastasis; precancerous lesion; rheumatoid arthritis; psoriasis; diabetic retinopathy; retinopathy of prematurity; macular degeneration; corneal graft rejection; neovascular glaucoma; retrolental fibroplasia; rubeosis; obler webber Syndrome; myocardial angiogenesis; plaque neovascularisation; telangiectasia; haemophiliac joint; angiofibroma; wound granulation; intestinal adhesion; atherosclerosis; scleroderma; hypertrophic scar.

Synthetic.

WO200253141-A2.

11-JUL-2002

14-DEC-2001; 2001WO-US048458.

(COLE-) COLEY PHARM GROUP INC 14-DEC-2000; 2000US-0255534P.

Bratzler RL;

involves administering at least one nucleic acid molecule to the subject. Inhibiting angiogenesis in a subject, WPI; 2002-566690/60. antiangiogenic

Claim 2; Page 37; 276pp; English.

The invention relates to inhibiting angiogenesis in a subject, comprising administering at least one antiangiogenic nucleic acid molecule. Also

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included is a kit comprising a first container housing the antiangiogenic mucleic acids, and instructions for administering them to a subject having a condition characterised by unwanted angiogenesis. The method is useful for inhibiting angiogenesis associated with solid tumour growth, tumour metastasis, precancerous lesion, rheumatoid arthritis, psoriasis, diabetic retinopathy, retinopathy of prematurity, macular degeneration, corneal graft rejection, neovascular glaucoma, retrolental fibroplasia, rubeosis, Osler-Webber Syndrome, myocardial angiogenesis, plaque neovascularisation, telangiectasia, haemophiliac joints, angiofibroma, who magnation and the present seminary and the present seminant and the present seminary and the present s
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  hypertrophic scars. The present sequence is an antiangiogenic nucleic
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Pred. No. 1e+03;
0; Mismatches 2; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1547 GCCTTCGGTCTTCGTCG 1563
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Human LSR gene biallelic marker upstream PCR primer #2.

(first entry)

21-MAY-2002

ABK41307;

BP.

ABK41307 standard; DNA; 20

.. 0

Gaps

.; 0

Human, obesity associated-biallelic marker, ss; LSR; USF2; PCR; primer, diquy response, hyperuriaemia; digestive pathology; hypertinsion; cancer; hepatic function disorder; cardiovascular disease; hyperlipidaemia; insulin disorder; atheromatous disease; cardiac insulficiency; obesity.

WO200206525-A2

24-JAN-2002.

28-JUN-2001; 2001WO-IB001477.

18-JUL-2000; 2000US-0219704P.

(GEST) GENSET.

Chumakov I, Abderrahim H, Bihain B; Cohen D, Blumenfeld M,

WPI; 2002-155043/20.

Set of novel map-related biallelic markers, preferably located on obesity disorder-associated chromosomal regions on chromosomes 3, 10 and 19, useful, for e.g. detecting statistical correlations between marker allele and a phenotype

Disclosure; Page 307; 311pp; English.

The invention relates to a set of novel map-related biallelic markers, preferably located on obesity disorder-associated chromosomal regions on chromosomes 3, 10 and 19. The markers are useful for genotyping or estimating the frequency of an allele in a population, for detecting an association between a genotype or haplotype and a phenotype, e.g. a disease involving drug responses, obesity or disorders related to obesity, such as hyperunicaemia, digestive pathology, hepatic function disorders, cancer, cardiovascular disease, hypertension, hyperlipidaemia, insulin disorders, atheromatous disease and cardiac insulficiency. The markers are useful for detecting a statistical correlation between a biallelic marker allele and a phenotype and/or between a biallelic marker

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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The invention relates to a map of the bread wheat D genome comprising the genome location of a microsatellite marker selected from a group of 185 such markers (ABQ921918-148092917). The invention also encompasses the use of left (ABQ921918-148092917). The invention also encompasses the use of left (ABQ921918-148092917). The invention also encompasses the use of left (ABQ92108-148093187) primers to amplify and detect the microsatellite markers, and to identify genes responsible for a phenotypic trait of interest in wheat. Wheat is an allohexaploid species consisting of 3 alploid encomes designated A, B and D, resulting from two successive intercrossings involving at least three different species. The D genome is thought to have been introduced in the cost recent intercrossing, between the amphiploid AABB and Triticum tauschii (DD), probably involving only a limited number of genotypes of both species. Due to its polyploid genome, the large size of its genome, and its low level of polymorphism, the genetic mapping of wheat has to between one and six nucleotides long, and are very polymorphic in length, mainly due to polymerase slippage during replication. This high degree of polymorphism makes them especially suitable for the genetic mapping of
                                                                                                                   ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Map of wheat D genome comprising the genome location of a microsatellite marker, useful for e.g. identifying genes responsible for a desired phenotypic trait, especially quantitative trait loci in wheat, and
haplotype and a phenotype. Sequences ABK41306-ABK41309 and ABK41328-ABK41331 represent PCR primers used to amplify human LSR gene or USF2 gene biallelic markers
                                                                                                                    Gaps
                                                                                                                                                                                                                                                                                                                                                 T. tauschii/wheat D genome microsatellite cfd60 right PCR primer.
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                                                                                                                                                                                                                                                                                                                                                                          Microsatellite marker; wheat; D genome; mapping; genotyping; polymorphism; phenotypic trait; QTL; quantitative trait locus; disease-associated gene; development factor; quality factor; resistance factor; wheat product; identification; detection; genetically modified wheat; PCR; primer; ss.
                                                                                   0.8%; Score 13.8; DB 1; Length 20; 88.2%; Pred. No. 1e+03; ative 0; Mismatches 2; Indels
                                                          Sequence 20 BP; 5 A; 4 C; 5 G; 6 T; 0 U; 0 Other;
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                                                                                                                                               1417 CGAAATCGGATCTCCGC 1433
                                                                                                                                                                                                                                               AB093162 standard; DNA; 20 BP.
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                                                                                                                   Matches 15; Conservative
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                                                                                        Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Aegilops tauschii.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           EP1217079-A1.
                                                                                                                                                                                                                                                                                                     29-AUG-2003
21-OCT-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        26-JUN-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Bernard M,
                                                                                                                                                                                                                                                                            ABQ93162;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           diseases.
                                                                                                                                                                                                                    RESULT 1516
                                                                                                                                                                                                                                  ABQ93162,
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Species which show little intraspecies polymorphism, such as wheat. In addition, microsatellites are codominant, and exhibit Mendelian continues and inheritance. The 185 microsatellite markers of the invention are developed from the ancestral diploid donor species Triticum tauschii and map to the wheat D genome, which is less polymorphic than the A or B compones. These microsatellite markers thus help to overcome some of the problems associated with the genetic mapping of wheat. The wheat D genome compones the microsatellite markers and associated primers of the invention are useful for identifying genes responsible for a phenotypic trait of interest, most notably QTDs (quantitative trait loci). In continuar they may be used for analysing genes and alleles implicated in conference and for identifying development factors, quality factors and disease and for identifying development factors, quality factors and cators conferring resistance to pathogens and xenobiotics. The mapping and genotyping diploid and polyphoid species of Triticum cum. Triticum comprises wheat or a related species; for identifying cultivars and hybrids of Triticum and related species; to assess whether or not a product comprises wheat or a related species; to assess whether or not a product comprises wheat or a related species; to assess whether or not a product comprises wheat or a related species; to assess whether or not a microsatellite marker right PCR primer of the invention. (Updated on 29-conference of the invention. (Updated on 29-conference of the invention.)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Indels
which show little intraspecies polymorphism,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 20 BP; 4 A; 3 C; 7 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          996 CCTGCTCATCAACGAGA 1012
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      08-MAR-2001; 2001US-0274191P.
22-MAR-2001; 2001US-0277826P.
29-MAR-2001; 2001US-0279840P.
13-ARR-2001; 2001US-0282981P.
13-ARR-2001; 2001US-0283656P.
31-JUL-2001; 2001US-0309247P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ABS51129 standard; DNA; 20 BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2000US-0258171P.
2001US-0269940P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Local Similarity
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20-FEB-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              21-DEC-2000;
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10017621-3sl.rng

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The present invention relates to the isolation of novel human proteins referred to as NOVX, and the polynucleotide sequences encoding them. The NOVX proteins of the invention include NOV1-NOV13. NOVX proteins, NOVX mucleic acids and antibodies are useful for treating or preventing a NOVX particularly humans. Such disorders include cardiomypathy, atherosclerosis, diabetes, cancer (e.g. adenocarcinoma, lymphoma, processing and metabolic pathways, disorders related to cell signal processing and metabolic pathways, disorders related to cell signal system (e.g. those induced by trauma, surgery and/or neoplastic disorders), acquired immunodeficiency syndrome (AIDS), inflammatory
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              disorders (e.g. asthma) obesity, anorexia, cancer-associated cachexia, neurodegenerative disorders (e.g. Alzheimer's disease, Parkinson's disease, immune disorders, graft versus host disease, Crohn's disease, multiple sclerosis, haemophilia, idiopathic thrombocytopenic purpura, and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Muscular; cytostatic; nootropic; neuroprotective; ophthalmological; antilipeamic; osteopathic; caspase 6; Rieger's syndrom; bone metabolism; ataxia telangiectasia; hyperproliferative disorder; cholesterol disorder; haematopoietic disorder; cancer; neurological; Alzheimer's disease;
                                                                                               Liu X, Spytek KA, Zerhusen B;
Lepley DM, Burgess CE, Shimkets R;
Li L, Casman SJ, Boldog F, Gorman L;
Rieger D, Edinger S, Gunther E, Millet I;
cdougall J, Smithson G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          infections diseases (e.g. bacterial, fungal, protozoal or viral infections). The polynucleotide sequences are also useful in gene therapy. The present sequence represents a real time quantitative (RTQ)-
                                                                                                                                                                                                                                                              New NOVX polypeptides and polynucleotides, useful in gene therapy, particularly for treating or preventing cardiomyopathy, atherosclerosis, diabetes, Crohn's disease, hemophilia or cancer in humans.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mouse caspase 6 antisense inhibition related oligo SEQ ID No 119.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DB 1; Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 20 BP; 10 A; 3 C; 5 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0.8%; Score 13.8; DB 1
88.2%; Pred. No. 1e+03;
ve 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PCR primer used in NOVX expression studies
                                                                                                                Patturajan M, Grosse WM, Lepley DM, Szekeres E, Vernet CAM, Li L, Casmal Gangolli EA, Pernandes E, Rieger D, Sciore P, Ellerman K, Macdougall J,
                                                                                                                                                                                                                                                                                                                                                Example 2; Page 254; 391pp; English
                                                                                                   Tchernev V, Liu X,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1240 TTCATCTTCCGTATCTT 1256
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ВР
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    apoptotic; mouse; murine; ds.
10-AUG-2001; 2001US-0311754P.
17-AUG-2001; 2001US-0313331P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              03-OCT-2001; 2001WO-US030871
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               88.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 18 TICATCTICCGCATITI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAL40400 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local Similarity 88.2
Matches 15; Conservative
                                                           (CURA-) CURAGEN CORP.
                                                                                                                                                                                                                       WPI; 2002-508801/54.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WO200229066-A1
                                                                                                   Alsobrook JP,
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Gaps

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The invention relates to an antisense oligonucleotide compound of 8 to 50 nucleotides in length that is targeted to a nucleic acid molecule encoding caspase 6, where the oligonucleotide specifically hybridises with and inhibits the expression of caspase 6. The oligonucleotide of the invention specifically hybridises to and inhibits expression of caspase 6 in cells or tissues. The oligonucleotides can be administered or cissues. The oligonucleotides can be administered or condition associated with caspase 6, such as Rieger's syndrome or ataxia telangiectasia, hyperproliferative disorder, a haematopoietic disorder, a bone metabolism or cholesterol disorder, a various types of cancer, neurological conditions such as Alzheimer's disease and other deregulated apoptotic pathological conditions. This polynucleotide sequence regulated apoptotic pathological conditions. This polynucleotide sequence regulated amouse caspase 6 oligonucleotide relating to the invention. NOTE: This phosphorothioate oligonucleotide sequence has 2'-MOE wings and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human; antisense; cytohesin-1; guanine nucleotide exchange protein; ARF; ADP ribosylation factor; inflammation; antiinflammatory; tumour;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
                                                                                                                                                      An antisense oligonucleotide of 8 to 50 nucleotides in length that
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human cytohesin-1 3' UTR antisense oligonucleotide, ISIS#111045.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            .
0
                                                                                                                                                                        inhibits caspase 6, is useful for treating Rieger's syndrome
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0.8%; Score 13.8; DB 1; Length 20; 88.2%; Pred. No. 1e+03; Ve 0; Mismatches 2; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Seguence 20 BP; 4 A; 7 C; 6 G; 3 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (ISIS-) ISIS PHARM INC.
(BOEH ) BOEHRINGER INGELHEIM PHARM INC.
                                                                                   Watt AT;
                                                                                                                                                                                                         Claim 3; Page 92; 141pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            211 CAGATAGGCCTGGATGA 227
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               13
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              04-OCT-2000; 2000US-00679299.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      30-OCT-2001; 2001WO-US047583.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       22-FEB-2001; 2001US-00791243.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match 0.8%;
Best Local Similarity 88.2%;
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                                                                                   Zhang H,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABS73952 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            15; Conservative
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                                                (ISIS-) ISIS PHARM INC.
                                                                                                                   WPI; 2002-471315/50
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                                                                                   3rown-Driver VL,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                cytostatic; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    36-SEP-2002.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          a deoxy gap
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RESULT 1519
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Matches
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New antisense oligonucleotide encoding human cytohesin-1, useful for

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                                                                                                               uncleobases targeted to a nucleic acid molecule encoding human cytohesin-
1, specifically hybridises with, and inhibits the expression of, human cytohesin-i, a guanine nucleotide exchange protein for ARF (ADP ribosylation factor). The antisense compound may be used in a pharmaceutical composition for inhibiting the expression of cytohesin-linhuman cells or tissues, and in treating a disease or condition associated with cytohesin-l by administering to the human the antisense compound e.g. tumour or inflammation. The antisense compound is also useful for diagnostics, therapeutics, prophylaxis and as research
preventing or treating a disease or condition associated with cytohesin-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The present invention describes a method of arraying genome clones. The method comprises: (a) clones of the genomic libraries contained in multiwell plates numbered for discrimination are mixed in each of the multiwell plates; (b) a primer designed based on the chromosome marker sequence is added to the mixture to carry out an amplification reaction; (c) a signal corresponding to the marker is detected from the resultant amplified product to specify the discrimination Nos. of the multiwell plates containing the clones having said marker sequence; (d) the order of the markers is changed so that the same discrimination Nos. succeed to the maximum in the specified discrimination Nos. to array the multiwell plates; (e) the clones in the multiwell plates of the specified discrimination Nos. are mixed respectively in each wells of longitudinal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  chromosome 1p36-35; chromosome 21q22.1; genetic analysis; genome;
                                                                                                                                                                                                                                                                                                    reagents and kits. The present sequence is an antisense oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
                                                                                                   comprising 8-30
                                                                                                                                                                                                                                                                                                                                                                                                                                          0
                                                                                                                                                                                                                                                                                                                                                                                             0.8%; Score 13.8; DB 1; Length 20;
88.2%; Pred. No. 1e+03;
tive 0; Mismatches 2; Indels
                                                                                                   new antisense compound,
                                                                                                                                                                                                                                                                                                                                                         Sequence 20 BP; 1 A; 11 C; 5 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human chromosome 1p36-35 PCR primer SEQ ID NO:752.
                    expression e.g. tumor or inflammation.
                                                            Example 15; Page 81; 107pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 4; Page 19; 528pp; Japanese.
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                                                                                                                                                                                                                                                                                                                      targeting human cytohesin-1
                                                                                                 The invention relates to a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  733 GCACCCTGCACCGCCAT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          4 GCGCCCTGCACCGCCCT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ABL43708 standard; DNA; 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
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Matches 15, Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PCR primer; ss.
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resultant cultures are amplified by using the above primer; (g) signals are detected from the amplified products; (h) the clones in the multiwell plates are specified from the detected result; and (i) the clones are reconstituted as the positions on the chromosome and arrayed. The microarray is useful for gene analysis. ABL42957 to ABL45322 represent
                                                                                                                                                                                                                                                                                                                                                                        Human; MEXK4 modulation; mitogen-activated protein kinase kinase 4; MTK1; MPSTK4; MAPSTK4; MAPKKK4; cytostatic; prophylaxis; immunological; Myperproliferative disorder; cancer; therapy; antieense; inflammatory; phosphorothioate backbone; ss.
                                                                 PCR primers for human chromosome 1p36-35 DNA, and ABL45323 to ABL45634 represent PCR primers for human chromosome 21q22.1, which are specifically claimed for use in the present invention
                                                                                                                                                                    Gaps
and lateral directions; (f) the mixed clones are cultured and the
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                                                                                                                                        Score 13.8; DB 1; Length 20;
Pred. No. 1e+03;
0; Mismatches 2; Indels
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'note= "2'-methoxyethyl nucleotides"
                                                                                                                                                                                                                                                                                                                                                   Human MEKK4 antisense oligonucleotide, ISIS #123107.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      note= "2'-methoxyethyl nucleotides"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     'note = "Phosphorothioate backbone"
                                                                                                                    Sequence 20 BP; 6 A; 7 C; 4 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Location/Qualifiers
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base= m5c
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AAD37172 standard; DNA; 20
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                                                                                                                                                                                                                                                       RESULT 1521
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Claim 36; SEQ ID NO 5188; 167pp + Sequence Listing; English
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                                                                                                                                                                                   The present invention relates to antisense compounds, compositions and methods for modulating the expression of MEKK4 (also referred as mitogenactivated protein kinase kinase 4; MAPIZK4; MAPIZKK kinase kinase 4; MAPIZKK, MTK1). The antisense oligos are useful for inhibiting the expression of MEKK4 in cells or tissues. They are also useful for treating an animal having a disease or condition associated with MEKK4 such as immunological, inflammatory, hyperproliferative disorder or cancer. Sequences of the invention are also useful for they are also useful in antisense therapy. The present sequence is an antisense oligomucleotide targetted to human MEKK4 DNA. This sequence is used in the exemplification of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human; methylated gene; methylation; breast cancer; marker; WT-1; cell proliferative disorder; TWIST; HOXAS; NES-1; RARbeta; cyclin D2; retinoic acid receptor beta; oestrogen receptor; Wilms' tumour; 14.3.3 sigma; HIN-1; RASSF1A; tumour suppressor gene; hypermethylation;
                                                                                                                                                                                                                                                                                                                                                                   Gaps
                                                                                                                         nucleic acid encoding mitogen-
for treating immunologic disorder,
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0; Mismatches
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                                                                                 Wyatt JR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Cyclin 14-3-3 sigma gene PCR primer #14.
                                                                                                                        New antisense compound targeted to activated protein kinase 4, useful inflammatory disorder or cancer.
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                                                                                                                                                                Claim 3; Page 92; 132pp; English
                                                                                 Monia BP,
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                    28-SEP-2001; 2001WO-US030549.
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                                         29-SEP-2000; 2000US-00676436
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Best Local Similarity 88.2
Matches 15; Conservative
                                                            (ISIS-) ISIS PHARM INC.
                                                                                Gaarde WA,
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                                                                                                    WPI; 2002-416486/44.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PCR; primer; ss
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04-APR-2002
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                                                                                Ward DT,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             colloid carcinoma, tubular carcinoma, medullary carcinoma, metaplastic acarcinoma, intraductal carcinoma in situ, lobular carcinoma in situ and papillary carcinoma in situ. The present sequence is a primer used in the exemplification of the invention
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Diagnosing and/or determining a predisposition to a cellular problemative disorder of breast tissue, in particular breast cancer, by determining the state of methylation of one or more nucleic acids isolated from the subject.
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signal transduction; DNA replication; cell division; growth;
proliferation; Candida albicans; fungicide; antifungal; PCR; primer; ss.
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                                                                                                                                                                                                 Claim 12; Page 46; 115pp; English.
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20-FEB-2001; 2001US-00792024.
22-AUG-2001; 2001US-0314050P.
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Matches 15; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Roemer T, Jiang B,
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The invention relates to constructing (M1) a strain of diploid fungal cells in which both alleles of a gene are modified, comprising modifying cone allele by insertion or replacement by a cassette having an expressible selectable marker and modifying other allele by insertion of a promoter replacement fragment with a heterologous promoter, so that expression of the second allele is regulated by the promoter. (M1) is useful for constructing a strain of diploid fungal cells in which both alleles modified are medified. The diploid fungal cells in which both alleles modified are useful for identifying a gene that contributes to the survival or growth of a fungus, a gene that contributes to the virulence and/or pathogenicity of a fungus, a gene that contributes to the resistance of a diploid fungus to an antifungal agent, an antifungal agent that inhibits the growth of a diploid fungus contributes to the resistance of a diploid fungus to an antifungal agent, an antifungal agent that inhibits the growth of a diploid fungus contributes to the resistance of a diploid fungus to an antifungal agent, an antifungal agent that inhibits the growth of a gene product, preferably enzymatic activity, carbon cativity of a gene product, preferably enzymatic activity, carbon transluction, preferably enzymatic activity, carbon cativity. The method is useful for identifying a compound having the ability to inhibit growth or proliferation of c. albicans cells and for treating infection by C. albicans. The present sequence is that of a PCR continuation is not represented in the printed sequence is that of a presented in the printed superinted superinted for the printed for a presented in the method of the invention. Note: The sequence data for this patent is not represented in the printed sequence is that of a presented in the method of the invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       this patent is not represented in the printed specification but is based on sequence information supplied to Derwent by the European Patent Office
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Pungus; yeast; tetracyclin; promoter; GRACE strain; biosynthesis;
signal transduction; DNA replication; cell division; growth;
proliferation; Candida albicans; fungicide; antifungal; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 20 BP; 3 A; 7 C; 5 G; 5 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1717 CTGAGCCATGTTCACCT 1733
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Boone C,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             4 CTGAGCCTTGTGCACCT 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ABZ31379 standard; DNA; 20 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       26-DEC-2001; 2001WO-US049486.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       29-DEC-2000; 2000US-0259128P.
20-FEB-2001; 2001US-00792024.
22-AUG-2001; 2001US-0314050P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           30-JAN-2003 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (ELIT-) ELITRA PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Roemer T, Jiang B,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2002-566694/60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Candida albicans
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WO200253728-A2.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ABZ31379;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        RESULT 1524
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The invention relates to constructing (M1) a strain of diploid fungal cells in which both alleles of a gene are modified, comprising modifying cells in which both alleles of a gene are modified, comprising modifying cone allele by insertion or replacement by a cassette having an expressible selectable marker and modifying other allele by recombination, of a promoter replacement fragment with a heterologous promoter. (M1) is useful for constructing a strain of diploid fungal cells in which both alleles of a gene are modified. The diploid fungal cells in which both alleles modified are useful for identifying a gene that contributes to the virulence and/or pathogenicity of a fungus, a gene that contributes to the resistance of a diploid fungus to an antifungal agent that inhibits the growth of a diploid fungus and for identifying a therapeutic agent for treatment of a mammalian disease. (M1) is useful for identifying a compound which modulates the activity of a gene product, preferably enzymatic activity, carbon compound catabolism, blosynthetic, transporter, transcriptional, cranslational, signal transduction, DNA repplication and cell division activity. The method is useful for identifying a compound having the contrainty of inhibit growth or proliferation of c. albicans cells and contrainty of a perent is not repersented in the printed specification but is based to contrain the printed specification but is based this patent is not represented in the printed specification but is besent of contraints.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       o;
                                                                                                                                                                                                                                                                                                                                                                                                                                                      is patent is not represented in the printed specification but is based sequence information supplied to Derwent by the European Patent Office
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CPRA, CPRB, CYP52A1A, CYP52A2A, CYP52A2B, CYP52A3A, CYP52A3B, CYP52A5A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CYPS2ASB; CYPS2ABA; CYPS2ABB; CYPS2D4A; URA3A; cytochrome P450; NADPH2 reductase; omega-hydroxylase complex; dicarboxylic acid; ss; quantitative competitive reverse transcription PCR; QC-RT-PCR; primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Candida tropicalis CYP52A5A/CYP52A5B gene QC-RT-PCR primer 7581-97-F.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0
Claim 36; SEQ ID NO 5598; 167pp + Sequence Listing; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0.8%; Score 13.8; DB 1; Length 20;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1335 AGCCGAGGCCCTTTTGA 1351
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98US-0103099P.
99US-0123555P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     88.2%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (revised)
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MADDURI K M.
CORNETT C A.
BRENNER A A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WILSON C R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CRAFT D L.
EIRICH L D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Best Local Similarity
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05-0CT-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ABK31851;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
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(MADD/)
(CORN/)
(BREN/)
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ABK31851
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                                                                                                                                                                        Adipose protein; ss; adp; obesity; transgenic animal; obesity; adipositas; bulimia; wasting; cachexia; eating disorder; body weight disorder; aneight loss; cancer; infectious disease; hypogonadism; Prader-Willi syndrome; Laurence-Moon-Biedl syndrome; hypothyroidism; diabetes; Cushing's syndrome; endocrine disorder; hypothyroidism; diabetes; inflammatory bowel disease; PCR primer; ulcerative colitis; anorexia nervosa; glycogen storage disease; lipid storage disease; lipoma.; liposarcoma; heart disease; hypertension; infertility; acquired immunodeficiency syndrome; AIDS.
                                                                                                     Novel isolated nucleic acid encoding cytochrome P450 and NADPH reductase enzymes of omega-hydroxylase complex of Candida tropicalis, useful for increasing production of dicarboxylic acids.
                                               Cornett CA;
                                                                                                                                                                                                                                                                                                                                                                               Gaps
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                                                                                                                                                                                                                                                                                                                                                    0.8%; Score 13.8; DB 1; Length 20;
88.2%; Pred. No. 1e+03;
tive 0; Mismatches 2; Indels
                                              Madduri KM,
                                                                                                                                                                                                                                                                                                                                Sequence 20 BP; 7 A; 3 C; 9 G; 1 T; 0 U; 0 Other;
                                              Eshoo M,
Gleeson M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mouse adipose protein, adp, PCR primer #4.
                                                                                                                                                    Example 11; Col 35-36; 173pp; English.
                                              Eirich LD,
                                                         Loper JC,
                                                                                                                                                                                                                                                                                                                                                                                                      1010 AGAGGGGAGAGCTCAAG 1026
                                                                                                                                                                                                                                                                                                                                                                                                                          2 AGAGGGCAGGGCTCAAG 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              16-JUN-2000; 2000US-0211914P.
23-JUN-2000; 2000EP-00113049.
28-JUN-2000; 2000US-0214518P.
17-APR-2001; 2001EP-00109537.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        13-JUN-2001; 2001WO-EP006713.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ABK16359 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
                                                                                                                                                                                                                                                                                                                                                                               Conservative
                                             Craft DL,
Tang M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (DEVE-) DEVELOGEN AG
                                                                                WPI; 2002-138383/18
        LOPER J C.
GLEESON M.
                                                                                                                                                                                                                                                                                                                                                                  Local Similarity
nes 15; Conserv
TANG M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WO200196371-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mus musculus.
                                                         Brenner AA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   14-MAR-2002
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                                              g,
                                                                                                                                                                                                                                                                                                                                                       Query Match
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           (LOPE/)
(GLEE/)
(TANG/)
                                              Wilson
                                                                                                                                                                                                                                                                                                                                                                    Best Loca
Matches
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polypeptide which regulates, causes or contributes to obesity in an animal or a human. The polymolectides, proteins, ant-adp antibodies, modulators of adp activity, adp antisense nucleic acids, expression vectors, adp transgenic animals are useful in the diagnosts and treatment of obesity, adipositas, bulimia, wasting (cachexia), eating disorders and/or disorders of body weight/body mass, weight loss due to cancer or infectious diseases, genetic disorders associated with hypogonadism e.g. prader-Willi syndrome, Laurence-Moon-Biedl syndrome, hypothyroidism, cliseases, inflammatory bowel disease, uncerative colitis, and anorexia nervosa. They are also useful for treating disorders of body weight/mass treating lipomas, and/or liposarcomas. The compositions are also useful for treating disorders of body weight/mass treating lipomas, and/or liposarcomas. The compositions are also useful for treating conditions associated with under weight e.g. enhancing or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     controlling fertility, controlling weight loss in acquired immunodeficiency syndrome (AIDS) or cancer patients. The present sequence is a PCR primer used to amplify an adp nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Raf kinase; hyperproliferation; neovascularisation; ocular angiogenesis; therapy; cancer; cytostatic; anti-angiogenic; vascular; ophthalmological;
                                                                                Novel nucleic acid encoding adipose polypeptide which regulates, causes or contributes to obesity, useful for treating obesity, heart disease, hypertension, infertility, and controlling weight loss in cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
                                                                                                                                                                                                          The invention relates to a nucleic acid encoding a adipose (ADP)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ó
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0.8%; Score 13.8; DB 1; Length 20;
86.2%; Pred. No. 1e+03;
ve 0; Mismatches 2; Indels
                Rothe M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human raf kinase related antisense oligonucleotide #17.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 20 BP; 6 A; 7 C; 4 G; 3 T; 0 U; 0 Other;
              Haeder T,
              Dohrmann C,
                                                                                                                                                                        Claim 1; Page 158; 188pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      883
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96US-00756806.
97US-00888982.
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98US-00143214
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      867 GCAGTACCTGGATGACT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GGAGTGCCTGGATGACT
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15; Conservative
              Ciossek T,
                                                 WPI; 2002-106464/14.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         antisense; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   13-DEC-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Unidentified
              Broenner G,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            JS6410518-B1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     31-MAY-1995;
26-NOV-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               25-JUN-2002.
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Best Local S
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(ISIS-) ISIS PHARM INC

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Best Local Similarity
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                                                                                                                                      interactions
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                         22-AUG-2002
                                                                                                                                                                         Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     07-MAY-2002.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                        ABQ66488;
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o
                                                                                                                                                                   The present invention relates to novel antisense oligonucleotides which are targetted to nucleic acids encoding human raf proteins and capable of inhibiting raf expression. The invention also relates to methods of inhibiting hyperproliferation of cells which involves contacting the hyperproliferating cells with a therapeutically effective amount of an oligonucleotide of the invention. The method is useful for treating cancer, angiogenesis or neovascularisation, especially ocular angiogenesis or neovascularisation. The present DNA sequence is human raf kinase related antisense oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The sequence represents a mouse syndecan-1 reverse transcription PCR primer. The invention relates to a novel use of a Smad3 inhibitor in preparing a medicament to treat or prevent wounds or fibrosis. The invention has antifibrotic and vulnerary activity. The Smad3 inhibitors are useful for preventing fibrosis and improving wound healing. The Smad3 protein, polypeptides and peptide fragments are useful for generating antibodies, as reagents for research purposes, or the identification of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         preventing wounds or fibrosis, or as reagents in assays for screening compounds for preventing fibrosis and improving of wound healing.
                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
                                                                                     cancer, angiogenesis or neovascularization by administering
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Smad3 inhibitors in preparing a medicament for treating or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Smad3; wound healing; fibrosis; antifibrotic; vulnerary; mouse; PCR primer; reverse transcription; syndecan-1; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                         .<del>.</del>
                                                                                                    oligonucleotides targeted to human raf sequences.
                                                                                                                                                                                                                                                                                                                                                                                   0.8%; Score 13.8; DB 1; Length 20; 88.2%; Pred. No. 1e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ວີ
                                                                                                                                                                                                                                                                                                                                                                                                                        Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mouse syndecan-1 reverse transcription PCR primer #2.
                                                                                                                                                                                                                                                                                                                                                    Sequence 20 BP; 6 A; 10 C; 0 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mitchell JB,
                                                                                                                                                                                                                                                                                                                                                                                                                        7
                                                                                                                                                                                                                                                                                                                                                                                                                        0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GS, Russo A,
                                                                                                                                        Disclosure; Col 59; 41pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example; Page 38; 65pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                            1168
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ABA96039 standard; DNA; 20 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             \vdash
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 19-MAY-2000; 2000WO-US013725
                                                                                                                                                                                                                                                                                                                                                                                                                                                          1152 TGACATGTGGGGTGTGG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           17 TGAGATGTGTGTGG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                         Matches 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Ashcroft
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2002-075348/10.
                                                    WPI; 2002-597918/64.
                                                                                                                                                                                                                                                                                                                                                                                                     Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WO200189556-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   19-MAY-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    08-APR-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                29-NOV-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Roberts AB,
                                                                                                         antisense
                  Monia BP;
                                                                                   Treating
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ABA96039
                                                                                                                                                                                                                                                                                                                                                                                       Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1528
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other cellular gene products involved in the regulation of fibrosis and improvement of wound healing, as reagents in assays for screening for compounds that can be used in the prevention of fibrosis and improvement of wound healing, and as pharmaceutical reagents in protecting against fibrosis and improving wound healing related to Smad3. Compounds that fibrosis and improving mound healing related to Smad3. Compounds that mutant Smad3 gene products, in inhibiting the activity of wild type and/or mutant Smad3 gene products, in elaborating the biological function of smad3, and in identifying compounds that disrupt normal Smad3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    as research and diagnostic reagents. Sequences ABQ66412-ABQ66511 represent chimeric phosphorothicate oligonucleotides, with 2'-MOB wings and a deoxy gap. The claimed sequences inhibit production of cytohesin-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           cytonesin-1; CT1; inhibit; cytostatic; antiinflammatory; cytostatic;
anti-infective; antisense gene therapy; infection; inflammation; tumour;
human; ss; inhibitor.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New antisense compounds directed against human cytohesin-1, useful treating and preventing infection, inflammation and tumors.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 13.8; DB 1; Length 20; Pred. No. 1e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                 Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      5:
                                                                                                                                                                                                                                                                                                                                                       4 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                             Score 13.8; DB 1;
Pred. No. 1e+03;
0; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human cytohesin-1 mRNA levels inhibitor #57.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 14; Col 41; 40pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1093 ACACTGTGGTACCGGCC 1109
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                         88.2%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1 ACACTGTGGAACCAGCC
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                                                                                                                                                                                                                                                                                                                                                           BP; 5 A; 7 C;
                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match 0.8
Best Local Similarity 88.2
Matches 15, Conservative
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Indels

Mismatches

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15; Conservative

Matches

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Gaps

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oligonucleotide probes (I) for use on a support to which complementary oligonucleotide probes (II) will hybridise with little mismatch, where (I) have melting temperatures within a narrow range. The method is useful for detecting infectious diseases caused by bacterial infectious agents e.g. Salmonella, Listeria monocytogenes and Haemophilus influenza, fungal infectious agents e.g. Cryptococcus neoformans, Candida albicans and Aspergillus fumigatus, viruses e.g. T-cell lymphocytotrophis cirus, Epstein-Barr virus and polio virus, and parasitic infectious agents medinesis. The method is also useful for detecting genetic diseases such as 21 hydroxylase deficiency, Turner Syndrome and obesity defects.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Detecting cancer involving oncogenes, tumour suppressor genes, or genes involved in DNA amplification, replication, recombination or repair, the cancer is specifically associated with a gene selected from BRCAI gene, p53 gene, human papillomavirus types 16 and 18 and liver cancers. The method is also used for environmental monitoring, forensics and the food and feed industry, detecting comprises scanning (using e.g. a scanning electron microscope and infrared microscope) the support at the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       oligonucleotide probe
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   particular sites and identifying if ligation of the oligonucleotide probe
sets occurred and correlating (using a computer) identified ligation to a
presence or absence of the target nucleotide sequences. ABI82074 to
ABI97546 represent oligonucleotide sequences used in the exemplification
                                                                                                                                                                                                                                                                                                                                   Human, K-ras, PCR primer, probe, capture probe, mutation detection, ligase detection reaction; LDR; p53; BRCA1; BRCA2; infectious disease; infection; 21 hydroxylase deficiency; Turner Syndrome; obesity; cancer; oncogene; tumour suppressor; human papillomavirus; forensic; environmental monitoring; food industry; feed industry; ss.
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Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 20 BP; 5 A; 8 C; 4 G; 3 T; 0 U; 0 Other;
2
                                                                                                                                                                                                                                                                                                  Capture oligonucleptide Zip ID#2505 oligo #9.
Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example 5; Fig 29; 300pp; English
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                                       749
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                                                                                                                                                                            ABI95418 standard; DNA; 20 BP
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                                     733 GCACCCTGCACCGCCAT
                                                                          GCGCCCTGCACCGCCCT
                                                                                                                                                                                                                                                          (first entry)
Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Barany F, Zirvi M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2002-034366/04
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WO200179548-A2.
                                                                                                                                                                                                                                                          16-FEB-2002
15;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                           Synthetic
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                                                                                                                                                                                                                   ABI95418;
                                                                                                                                    RESULT 1530
Matches
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oligonuclectide probes (I) for use on a support to which complementary oligonuclectide probes (II) will hybridise with little mismatch, where (I) have melting temperatures within a narrow range. The method is useful for detecting infectious diseases caused by bacterial infectious agents e.g. Salmonella, Listeria monocytogenes and Haemophilus influenza, fungal infectious agents e.g. Cryptococcus neoformans, Candida albicans and Aspergillus fundgautus, viruses e.g. T-cell lywphocytotrophis cirus, Epstein-Barr virus and polio virus, and parasitic infectious agents selected from Onchoverva volvulus, Entamobal histolytica and Dracunculus medinesis. The method is also useful for detecting genetic diseases such as 21 hydroxylase deficiency, Turner Syndrome and obesity defects.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Detecting cancer involving oncogenes, tumour suppressor genes, or genes involved in DNA amplification, replication, recombination or repair, the cancer is specifically associated with a gene selected from BRCAH gene, p53 gene, human papillomavinus types 16 and 18 and liver cancers. The method is also used for environmental monitoring, forensics and the food and feed industry, detecting comprises scanning (using e.g. a scanning electron microscope and infrared microscope) the support at the particular sites and identifying if ligation of the oligonucleotide probe
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   sets occurred and correlating (using a computer) identified ligation to a presence or absence of the target nucleotide sequences. ABI82074 to ABI97546 represent oligonucleotide sequences used in the exemplification
                                                                                                                                                                                                                                                                                                                                                                      Human, K-ras, PCR primer, probe, capture probe, mutation detection,
ligase detection reaction, LDR, p53, BRCA1; BRCA2, infectious disease,
infection, 21 hydroxylase deficiency, Turner Syndrome, obesity, cancer,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Designing capture oligonucleotide probes for use on a support to which complementary oligonucleotides hybridize with little mismatch.
Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The present invention describes a method (M1) for designing capture
                                                                                                                                                                                                                                                                                                                                                                                                                                        oncogene; tumour suppressor; human papillomavirus; forensic; environmental monitoring; food industry; feed industry; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Kliman R;
  2,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 20 BP; 5 A; 5 C; 8 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                             Capture oligonucleptide Zip ID#518 oligo #9.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Favis R,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 5; Fig 29; 300pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gerry NP,
                                         288 ACTICGITCIGCACGGG 304
                                                                                                                                                                                               ABI93431 standard; DNA; 20 BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          04-APR-2001; 2001WO-US010958.
                                                                                    18 AGTICGITCGGCACGGG
                                                                                                                                                                                                                                                                                    (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        of the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Zirvi M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2002-034366/04.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Synthetic.
                                                                                                                                                                                                                                        ABI93431;
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Score 13.8; DB 1; Length 20; Pred. No. 1e+03;

0.8%;

Query Match Best Local Similarity

DB 1; Length 20;

Score 13.8; DB Pred. No. 1e+03;

0.8%;

Best Local Similarity

Query Match

SS.

oligonucleotide;

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Matches

RESULT 1532

ABL50712

ABL50712;

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congretes Alloysons represent proportion considers and seek designed to target different regions of human talin RNA, and were analysed for their effect on talin expression. The antisense were calculated to target different regions of human talin RNA, and were analysed for their effect on talin expression. The and were calculated by the control of their editions of their editions of the calculation of callular and vinculin to integrins, thereby linking the regulation of callular and may play a role in platelet adhesion as its subcellular distribution differs between resting non-adhesive platelets subcellular distribution differs between resting non-adhesive platelets. The contractile force to the extracellular and activated adhesive platelets. It could also play a major role in determining muscle strength and cardiac function as it has been found to participate in the transmission of contractile force to the extracellular expression at myotendinous junctions. The expression of talin is downregulated by androgens in prostate tissues, a phenomenon known to contribute to the development of prostate cancer. The oligonucleotides of the invention are useful for diagnosis, prevention and treatment of talin the related disorders, such as those related to cellular adjection. The contraction and treatment of talin and the contraction and contraction and calculation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New antisense oligonucleotides targeted to nucleic acid molecules encoding human Talin, useful for inhibiting the expression of human Talin and for treating a human having a disease or condition associated with
                                                                                                                                                                                                                                                                                                                                                                                           /note= "This oligonucleotide has a phosphorothioate backbone and 2-"methyoxyethyl (2-'MOS) wings at the 5' and 3' ends, which are 5 nucleotides in length. Also all cytosine nucleotides are 5-methylcytosines"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          present sequence represents a human c-Ha-ras phosphorothioate antisense
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequences ADG90460-ADG90539 represent phosphorothioate targeted to the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  used as a positive control in determining optimal concentration for a particular cell line.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0.8%; Score 13.8; DB 1; Length 20; 88.2%; Pred. No. 1e+03; ve 0; Mismatches 2; Indels
                       cardiomyocyte; platelet; prostate; androgen downregulation; prostate cancer; talin-related disorder; adhesion-related disorder; expression inhibition; antisones there is the contract of the 
muscle strength; cardiac
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Cowsert LM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 20 BP; 5 A; 7 C; 4 G; 4 T; 0 U; 0 Other;
                                                                                                                                    antisense therapy; phosphorothioate; antisense
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         oligonucleotide used as a positive control
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Kishimoto TK,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ISIS PHARM INC.
BOEHRINGER INGELHEIM PHARM INC.
                                                                                                                                                                                                                                                             Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1571 ACTCAGGCAGGCCAGCT 1587
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                                                                                                                                                                                                                                                                                                                                                                   base
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Les 15; Conserv
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                                                                                                                                                                                                                                                                                            modified base
                                                                                                                                                                                                       Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             coupled receptor (GPCR) proteins. The GPCR sequences have antidiarrheic and laxative activities. The GPCR sequences can be used for developing drugs to treat diseases of the digestive organs, e.g. colitis, diarrhea, constipation and malabsorption syndrome, including gene diagnosis and therapy. The present sequence represents a PCR primer for rat GPCR, which
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human and rat brain-originated G protein-coupled receptor proteins and encoded DNAs, for developing drugs to treat diseases of the digestive organs, e.g. colitis, diarrhea, constipation and mal-absorption syndrome.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      present invention describes human and rat brain-originated G protein-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Rat; rZAQl; rZAQ2; G protein-coupled receptor; GPCR; antidiarrheic; laxative; drug development; digestive organ disease; colitis; diarrhoea; constipation; malabsorption syndrome; diagnosis; gene therapy;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
          Gaps
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          Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0;
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                                                                          567 CCTCCGTCGTGTCAGCC
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Terao Y,

Rattus sp.

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Gaps

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862

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Query Match

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Matches

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1533

RESULT

Human; mouse; ss; primer; gene 216; antiasthmatic; antiinflammatory; anorectic; chromosome 20pil-pl2; single nucleotide polymorphism; SNP; gene therapy; respiratory disease; asthma; obesity; PCR; bronchial hyper-responsiveness; chronic obstructive pulmonary disease; adult respiratory distress syndrome; inflammatory bowel syndrome.

Human gene 216 polymorphism detection PCR primer #32.

(first entry)

25-MAR-2003

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(ATP) binding cassette transporter gene subfamily C12, i.e. cystic bibrosis transmembrane conductance regulator/multidrug resistance-like subgroup (CFTR/MRP) family) polypeptide and its encoding polymucleotides The polypeptide is useful for screening agonists and antagonist of the ABCC12 polypeptide. The products of the invention are useful for screening an active ingredient for preventing and treating paraxysmal kinesigenic chorecathetosis or parhologies linked to dysfunction of transport of organic anion transporters such as cysteinyl leukotriene, anionic drugs, such as methotrexate, neutral drugs conjugated to acidic ligands, such as glutathione, glucuronate or sulphate conjugated drugs and can be used for somatic gene therapy. This sequence represents a region corresponding to an exon/intron boundary from the gene encoding a human ABCC12 isoform described in the disclosure of the invention
                                                                                                                                                                                         Adenosine triphosphate (ATP)-binding cassette transporter subfamily Cl cystic fibrosis transmembrane conductance regulator; human; CFTR/MRP; multidrug resistance-like subgroup; somatic gene therapy; ABCCL2; paroxysmal kinesigenic choreoathetosis; cysteinyl leukotriene; anionic drug; methotrexate; neutral drug; glutathione; glucuronate; sulphate conjugated drug; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New ATP-binding cassette transporter gene subfamily C12, ABCC12 polypeptide, useful for preventing or treating paroxysmal kinesigenic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      This invention describes a novel human ABCC12 (adenosine triphosphate
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                                                                                                                                                         Human ABCC12 exon 22/intron 22 boundary.
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HEALTH & HUMAN SERVICES
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                                        ABQ77206 standard; DNA; 20 BP
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                                                                                                                  (first entry)
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(USSH ) US DEPT
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Local St.
15;
                                                                                                                  24-APR-2003
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                                                                              ABQ77206;
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RESULT 1534
                                                                                                                                                                                                                                                                                                                                  Homo
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Dean M;

New isolated gene 216 nucleic acids, useful for diagnosing, preventing or treating a disorder, such as asthma, bronchial hyper-responsiveness, chronic obstructive pulmonary disease, obesity or inflammatory bowel

Example 10; Page 155; 650pp; English.

syndrome

Del Mastro RG;

Dupuis J,

Van Eerdewegh P,

Allen K, Pandit S;

WPI; 2003-092960/08.

Little RD,

Keith T, Simon J,

(SCHE) SCHERING CORP. (GENO-) GENOME THERAPEUTICS CORP.

13-APR-2001; 2001US-00834597. 13-APR-2001; 2001WO-US012245.

15-APR-2002; 2002WO-US012063.

WO200283077-A2

24-OCT-2002

Homo sapiens

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This invention relates to a novel isolated nucleic acid, gene 216, identified from human chromosome 20p13-p12. The invention also discloses regions of the 216 gene that contain single nucleotide polymorphisms (SNP's) which may be used as markers for disease susceptibility or severity. The nucleotides of the invention may have antiasthmatic, antinflammatory or anorectic activities and may be used in gene therapy. The nucleic acids, antibodies or its fragments are useful for diagnosing, preventing or treating a disorder, such as respiratory diseases (e.g. asthma, bronchial hyper-responsiveness, chronic obstructive pulmonary disease or adult respiratory distress syndrome), obseity, or inflammatory bowel syndrome. The nucleic acids are also useful for identifying increased susceptibility of a subject to the disorders mentioned. The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   nucleic acids can also be used as primers and templates for the recombinant production of disorder-associated peptides or polypeptides, for chromosome and gene mapping, or for tissue distribution studies. The present sequence represents a gene 216 specific PCR primer used in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
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Pred. No. 1e+03;
0; Mismatches 2; Indels
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Best Local
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1D ABX79
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DT 25-M2
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Gaps ..

881

865 AAGCAGTACCTGGATGA AGGCATTACCTGGATGA

Conservative

Matches

Similarity

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ABX74975 standard; DNA; 20 BP

RESULT 1535

ABX74975

ABX74975
ID ABX7
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AC ABX7

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The invention relates to antisense compounds targetted to a nucleic acid molecule encoding fibroblast growth factor (FGP) receptor 3 (also known as FGPR-3) ACH, TYR4 and CEK2) to inhibit its expression. Antisense compounds of the invention are useful for treating diseases or conditions associated with FGFR-3 such as developmental disorders or hyperproliferative disorders, especially cancer of colorectal, bladder, bone, lung, cervical, breast or skin. They are useful as research reagents, therapeutics, prophylaxis, kits and diagnostics, and as tools in differential and/or combinatorial analyses to elucidate expression patterns of a portion of the genes expressed within cells and tissues. They are also useful in antisense therapy. The present sequence is an antisense oligonucleotide targetted to human FGFR-3
                                                                                                                                                                                                                               /mod_base= OTHER /note= "Phosphorothioate backbone; All cytidine residues are 5-methylcytidines"
Human; antisense; fibroblast growth factor receptor 3; prophylaxis;
developmental disorder; hyperproliferative disorder; antisense therapy;
FGFR-3; ACH; JTK4; CEK2; cancer; phosphorothioate; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Novel compound targeted to a nucleic acid molecule encoding fibroblast
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               growth factor receptor 3, useful for inhibiting the expression of the receptor and for treating an animal having cancer or developmental
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0.8%; Score 13.8; DB 1; Length 20; 88.2%; Pred. No. 1e+03; 1ive 0; Mismatches 2; Indels
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                                                                                                                                                                                                                                                                                                                                                                /note= "2'-methoxyethyl (2'-MOE)
                                                                                                                                                                                                                                                                                                                                                                                                                                                       "2 -methoxyethyl (2'-MOE)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example 15; Page 79; 120pp; English.
                                                                                                                                                             Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                             OTHER
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                                                                                                                   Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Monia BP,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              disorder.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ACF57208;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 RESULT 1538
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Matches
                                                                                                  Homo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ACF57208/
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New isolated gene 216 nucleic acids, useful for diagnosing, preventing or treating a disorder, such as asthma, bronchial hyper-responsiveness, chronic obstructive pulmonary disease, obesity or inflammatory bowel
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                identified from human chromosome 20p13-p12. The invention also discloses regions of the 216 gene that contain single mucleotide polymorphisms (SNP's) which may be used as markers for disease susceptibility or severity. The nucleotides of the invention may have antiasthmatic, antinflammatory or anorectic activities and may be used in gene therapy. The nucleic acids, antibodies or its fragments are useful for diagnosing, preventing or treating a disorder, such as respiratory diseases (e.g. asthma, bronchial hyper-responsiveness, chronic obstructive pulmonary
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        inflammatory
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     nucleic acids can also be used as primers and templates for the recombinant production of disorder-associated peptides or polypeptides, for chromosome and gene mapping, or for tissue distribution studies. The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         sequence represents a gene 216 specific PCR primer used in the
                                               Human; mouse; ss; primer; gene 216; antiasthmatic; antinflammatory; anorectic; chromosome 20p13-p12; single nucleotide polymorphism; SNP; gene therapy; respiratory disease; asthma; obesity; PCR; bronchial hyper-responsiveness; chronic obstructive pullmonary disease; adult respiratory distress syndrome; inflammatory bowel syndrome.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     disease or adult respiratory distress syndrome), obesity, or inflammat
bowel syndrome. The nucleic acids are also useful for identifying
increased susceptibility of a subject to the disorders mentioned. The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    This invention relates to a novel isolated nucleic acid, gene 216,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Del Mastro RG;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human FGFR-3 antisense oligonucleotide, ISIS #125180.
         Human gene 216 polymorphism detection PCR primer #92.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Dupuis J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 20 BP; 4 A; 4 C; 7 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Ъ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example 10; Page 155; 650pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Van Eerdewegh
                                                                                                                                                                                                                                                                                                                                                                                                                                                              GENOME THERAPEUTICS CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  .
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13-APR-2001; 2001WO-US012245.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Little RD, Van Ees
Allen K, Pandit S,
                                                                                                                                                                                                                                                                                                                             15-APR-2002; 2002WO-US012063
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                scope of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                           SCHERING CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2003-092960/08
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity
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nucleotides"

Keith T, Simon J,

(SCHE) (GENO-) (

Homo

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Gaps

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Matches

RESULT 1537

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AAD55476;

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The present sequence is that of an antisense primer for Fas-associated ACR05705 generates a 695 bp product. Semi-quantitative RT-PCR was used to determine levels of FADD RNA in thyroids of gep transgenic mice during various stages of tumour development. These mice provide models of buman thyroid tumours. RADD expression was shown to decrease, in some cases to zero, during tumour development. It therefore provides a marker for the absence of in vivo tumour. FADD proteins are secreted from tumour cells. A low cellular amount and a high extracellular amount of FADD proteins are prognostic of resistance to chmotherapy. The invention provides methods for determining a status of tumour absence/presence, and for prognosis of the resistance of a tumour to chemotherapy on the basis of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Treating non-allergic inflammatory diseases, such as psoriasis, eczema, allergic contact dermatitis, latex dermatitis or inflammatory bowel
                                                                                                                                                                   Use of Fas associated protein with dead domain, and cellular phosphorylated p38-mitogen activated protein kinases as a biological
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Immunostimulatory, antiinflammatory, dermatological, antipsoriatic, antiuloer; gene therapy, vaccine, non-allergic inflammatory disease, psoriasis, eczema, allergic contact dermatitis, latex dermatitis, inflammatory bowel disease, ulcerative colitis, Crohn's disease, ss.
                                                                                        Buzyn A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0.8%; Score 13.8; DB 1; Length 20; 88.2%; Pred. No. 1e+03; ive 0; Mismatches 2; Indels
                                                                                        Ľ,
                                                                                        Michiels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 20 BP; 7 A; 7 C; 4 G; 2 T; 0 U; 0 Other;
                       (INRM ) INSERM INST NAT SANTE & RECH MEDICALE. (CNRS ) CNRS CENT NAT RECH SCI.
                                                                                        Feunteun J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Immunostimulatory nucleic acid #972.
                                                                                                                                                                                                                                                   Example 1; Page 50; 118pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1656 CCACACCCCTCACAGGG 1672
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        29-MAR-2002; 2002US-00112653.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ccacagrecreacaggg
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ACH03337 standard; DNA; 20
                                                                                      Tourneur L,
                                                                                                                                                                                                           indicator of tumor status.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first entry)
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                                                                                                                            WPI; 2003-645962/61.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Local Similarity
les 15; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Berg DJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         KRIEG A M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (KRIE/) KRIEG A M
(BERG/) BERG D J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        US2003050268-A1
                                                                                      Chiocchia G,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                25-SEP-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Krieg AM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ACH03337;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
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Matches
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        음
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ·
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The present invention describes a method and a kit for determining the expression of mRNA or CDNA of a protein participating in the maintenance of skin structure. The method is quantitative, simple and accurate in the determination of extracellular matrix components of laminin 5 chain genes LAMA3, LAMB3 and LAMC2, matrix metalloproteinases MMP-1, MMP-2, MMP-3 and MMP-9, VII collagen, type I collagen alpha 1 chain, type IV collagen alpha 2 chain, type IV collagen alpha 2 chain, TIMP-1, TIMP-2 and TIMP-3. ACE57201 to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FADD; mouse; tumour; marker; diagnosis; prognosis; thyroid; PCR; primer;
                                                                                Human, mouse, skin structure, skin, laminin 5 chain gene, LAMA3; LAMB3;
LAMC2; extracellular matrix component, matrix metalloproteinase, MMP-1,
MMP-2; MMP-3; MMP-9; TIMP-1, TIMP-2; TIMP-3; collagen, PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A method and a kit for determination of expression of mRNA or cDNA of protein participating in the maintenance of skin structure.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 13.8; DB 1; Length 20;
Pred. No. 1e+03;
); Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 20 BP; 9 A; 7 C; 4 G; 0 T; 0 U; 0 Other;
                                          reverse PCR primer SEQ ID NO:8.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 1; Page 2; 34pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     .,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1113 TGACATCCTGCTTGGGT 1129
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       BP.
                                                                                                                                                                                                                                                                                                          15-JAN-2002; 2002JP-00006797.
                                                                                                                                                                                                                                                                                                                                                15-JAN-2001; 2001JP-00006952.
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22-OCT-2002; 2002EP-00292619.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      23-DEC-2002; 2002WO-EP014906
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16-OCT-2003 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FADD antisense PCR primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                      (SHIS ) SHISEIDO CO LID
                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2003-407328/39
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Best Local Similarity
Matches 15; Conserv
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                                                                                                                                                                                                                        JP2002330792-A.
                                                                                                                                                                Homo sapiens
                                          Human LAMA3
                                                                                                                                                                                                                                                                  19-NOV-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   06-NOV-2003
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                                                                                                                                                                                 Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Treating and/or preventing allergy or asthma using an immunostimulatory nucleic acid alone or in combination with an asthma/allergy medicament.
                                                                              disease comprising administering to a subject having or at risk of developing a non-allergic inflammatory disease an immunostimulatory nucleic acid for prevention or treatment of the disease. The method is useful for treating non-allergic inflammatory diseases, such as psoriasis, eczem, allergic contact dermatitis, latex dermatitis or inflammatory bowel disease e.g., ulcerative colitis or Crohn's disease. This sequence represents an immunostimulatory nucleic acid
                                                                The invention describes a method of treating non-allergic inflammatory
                                                                                                                                                                                                                                                                                   Gaps
                                                                                                                                                                                                                                                                                     ;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ds; allergy; asthma; poly-G nucleic acid; aerosol formulation; hypo-responsive subject; immunostimulatory.
disease by administering an immunostimulatory nucleic acid.
                                                                                                                                                                                                                                                  DB 1; Length 20,
                                                                                                                                                                                                                                                                                   2; Indels
                                                                                                                                                                                                                   Sequence 20 BP; 2 A; 5 C; 6 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 20 BP; 2 A; 5 C; 6 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                  Score 13.8; DB 1
Pred. No. 1e+03;
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Fouron Y;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Immunostimulatory nucleic acid #929.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Disclosure; Page 19; 221pp; English.
                                 Disclosure; Page 35; 229pp; English
                                                                                                                                                                                                                                                                                     .,0
                                                                                                                                                                                                                                                                                                                     1547 GCCTTCGGTCTTCGTCG 1563
                                                                                                                                                                                                                                                                                                                                                    1 GCCTTCGATCTTCGTTG 17
                                                                                                                                                                                                                                                                                                                                                                                                                                        BP.
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                                                                                                                                                                                                                                                    0.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                          ADB37315 standard; DNA; 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first entry)
                                                                                                                                                                                                                                    Query Match
Best Local Similarity 88.2.
Thes 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           BRATZLER R L.
PETERSEN D M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2003-657977/62
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (PETE/) PETERSEN (FOUR/) FOURON Y.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           US2003087848-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         04-DEC-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADB37315;
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                                                                                                                                                                                                                                                                                                                                                                                                        RESULT 1541
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The invention relates to a compound 8-50 nucleobases in length targeted to a nucleic acid molecule encoding complement component C3. The compound peculically whybridises with the nucleic acid molecule encoding complement component C3 and inhibite the expression of complement component C3, or specifically hybridises with at least an 8-nucleobase component C3, an active site on a nucleic acid molecule encoding complement component C3. Also included are a composition comprising the compound and a paramaceutical carrier or diluent, inhibiting the expression of a pharmaceutical carrier or diluent, inhibiting the expression of component C3 in cells or tissues (comprising contacting the compound cited above) and treating an animal having a disease or condition associated with complement component C3 comprising administering to the animal the compound cited above so that expression of complement component C3 is inhibited. The antisense compounds are useful for inhibiting the expression of complement c3 in cells or tissues, or for treating an animal having a
                                                                                                                                                                                                                                                                                                                                                                                                                                           /note= "Phosphorothioate backbone and all cytosines are
                                                                                                                                                                                                                     Antinsense oligonucleotide targeting mouse C3 component, ISIS140104
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New antisense oligonucleotides targeted to a nucleic acid molecule encoding complement component C3, useful for treating a disease or condition associated with complement component C3, e.g. autoimmune
                                                                                                                                                                                                                                                   Mouse; ss; antisense; complement component C3; inflammation; septic shock; multiple organ failure; hyperacute organ failure; autoimmune disorder; CNS inflammation; multiple sclerosis; atherosclerosis; tumour.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /note= "2'-methoxyethyl nucleotides"
16. .20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               not\bar{e}= "2'-methoxyethyl nucleotides"
                                                                                                                                                                                                                                                                                                                                                                         Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example 16; Page 27; 72pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                            -methyl cytosines"
                                                                                                                                                                                                                                                                                                                                                                                                                           base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /mod_base= OTHER
1547 GCCTTCGGTCTTCGTCG 1563
                                 17
                                                                                                                      BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    23-OCT-2001; 2001US-00001076.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     23-OCT-2001; 2001US-00001076.
                                   1 GCCTTCGATCTTCGTTG
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                                                                                                                                                                                       (first entry)
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                                                                                                                    ADB90016 standard; DNA;
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                                                                                                                                                                                                                                                                                                                                                                                                             *tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   '*tag≈
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           disorder or infection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /mod
                                                                                                                                                                                                                                                                                                                                                                                                                                mod/
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Graham MJ, Watt AT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2003-606441/57
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    US2003096775-A1.
                                                                                                                                                                                                                                                                                                                                                                             Key
modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               modified_base
                                                                                                                                                                                                                                                                                                                                           Mus musculus
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                                                                                                                                                       ADB90016;
                                                                                      RESULT 1542
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Gaps

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Indels

7;

0.8%; Score 13.8; DB 1; Length 20; 88.2%; Pred. No. 1e+03;

Pred. No. 1e+03; 0; Mismatches

Conservative

Best Local Similarity Matches 15; Conserv

Query Match

'n

.; 0

Gaps

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autoimmune disorder (e.g. multiple sclerosis), an infection, or atherosclerosis, inflammation, septic shock, multiple organ failure, hyperacute organ failure and CNS inflammation. The compounds are also useful as research reagents and diagnostics, in distinguishing functions of various members of a biological pathway, or for preventing or delaying infection, inflammation or tumour formation. The present sequence is an antisense oligonucleotide targeting mouse C3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New compound, having a sequence targeted to a nucleic acid encoding human collapsin response mediator protein 2, useful for preparing a composition for treating hypercholesterolemia or hyperproliferative disorder, e.g.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /mod_base= OTHER
/note= "OTHER= phosphorothioate backbone, where 1-5 and
16-20 are 2' methoxyethyl nucleotides. All cytidines are
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                This invention relates to novel antisense oligonucleotides that modulate the expression of human eukaryotic translation initiation factor 2C 1 [EFF2C1]. EFF2C1 is located on chromosome 1p34-35, and is also known as Co-eFF2C, eFF2C, Golgi ER protein 95kDa, GER995 and Q99. It is an intracellular membrane associated protein thought to be involved in cellular differentiation, such that altered expression of EFF2C1 can
disease or condition associated with complement component C3 such as
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           antisense; ss; human; eukaryotic translation initiation factor 2C 1; EIF2C1, CO-eIF2C; eIF2C; Golgi ER protein 95kDa; GERp95; Q99; gene therapy; hyperproliferative disorder; familial hypercholesterolaemia; cancer; polycystic kidney disease; cystic fibrosis; progeriod syndrome; cytostatic; antilipaemic.
                                                                                                                                                                                                                                ٠,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Antisense oligo (SeqID 29) used to inhibit human EIF2C1 DNA.
                                                                                                                                                                                          Score 13.8; DB 1; Length 20;
Pred. No. 1e+03;
0; Mismatches 2; Indels
                                                                                                                                                                                                                              Indels
                                                                                                                                                                                                                              2;
                                                                                                                                                       Sequence 20 BP; 3 A; 9 C; 3 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 3; Page 76; 120pp; English.
                                                                                                                                                                                                                                                               407
                                                                                                                                                                                                                                                                                                                                                                                       BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  04-NOV-2002; 2002WO-US035324.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    08-NOV-2001; 2001US-00007078.
                                                                                                                                                                                             0.8%;
                                                                                                                                                                                                            88.2%;
                                                                                                                                                                                                                                                                                              20 rcadargaggracaggc
                                                                                                                                                                                                                                                             TCGGATGAGGTGCAGTC
                                                                                                                                                                                                                                                                                                                                                                                     ADB81512 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                         04-DEC-2003 (first entry)
                                                                                                                                                                                      Query Match 0.8 Best Local Similarity 88.2 Matches 15; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         . 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2003-449448/42.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Watt AT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WO2003040321-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      modified base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 L5-MAY-2003
                                                                                                                                                                                                                                                               391
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Ward DT,
                                                                                                                                                                                                                                                                                                                                               RESULT 1543
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affect cell growth, morphology and tumourigenicity. Accordingly, antisense oligonucleotides that inhibit the expression of BIF2C1 in cells or tissues can be used in gene therapy to treat various conditions including hyperproliferative disorders, familial hypercholesterolaemia and cancer, as well as polycystic kidney disease, cystic fibrosis and progeriod syndrome. As such, the oligos of the present invention can be described as having cytostatic and antilipaemic activities. This oligonucleotide sequence is an antisense oligo used to inhibit expression of the human eukaryotic translation initiation factor 2C I (BIF2C1) DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ò
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human; ss; PCR; retinal pigment epithelial-derived neurotrophic factor; PEDF; tumour; ocular disease; neuronal cell pathology; serine protesse; blood coagulation; thrombosis; bacterial infection; parasitic infection; elastosis; vascular disorder; fibrinoid formation; coagulation disorder; settic scalerosis; ischaemia; arthroses diabetes; emphysema; arthritis; septic shock; lung disease; complement activation; ulcerr; ulcerative colitis; pancreatitis; psoriasis; fibrinolytic disease; arthropathy; bone resorption; hypertension; congestive heart failure;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New purified retinal pigmented epithelium derived neurotrophic factor composition, useful for treating tumors, i.e. retinal tumor, ocular disease, neuronal cell pathologies, coagulation disorders or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human retinal pigment epithelial-derived factor (PEDF) PCR primer #1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Johnson LV;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         cirrhosis; protease allergy; chromosome 17p13.1-pter; primer.
                                                                                                                                                                                                                                                                                                                                                                                                       Score 13.8; DB 1; Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                    5;
                                                                                                                                                                                                                                                                                                                                           Sequence 20 BP; 3 A; 8 C; 2 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                 Pred. No. 1e+03;
0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 48; SEQ ID NO 9; 58pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             584
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92US-00952796.
95US-00520373.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADB99096 standard; DNA; 20 BP.
                                                                                                                                                                                                                                                                                                                                                                                                       0.88;
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                                                                                                                                                                                                                                                                                                                                                                                                                             88.2%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          568 CTCCGTCGTGTCAGCCT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                 Local Similarity 88.2
les 15; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (TOMB/) TOMBRAN-TINK J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STEELE F R.
CHADER G J.
BECERRA S P.
JOHNSON L V.
RODRIGUEZ I R
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2003-743982/70.
                                                                                                                                                                                                                                              of the human euka of the invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               US2003096750-A1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              24-SEP-1992;
29-AUG-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     04-DEC-2003
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                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
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(CHAD/)
(BECE/)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               RESULT 1544
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Matches
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      XXXX H H H X X B B B B B B B B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B
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Example 15; SEQ ID NO 71; 141pp; English.

gastric cancer.

colon or

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                                                                                                                                                                                                                                                                                                                                transforming growth factor beta receptor II; TGF-beta receptor II; hyperproliferative disorder; breast cancer; autoimmune disorder; rheumatoid arthritis; 2'-0-methoxyethyl gapmer;
                                                                                                                                                                                                                                                                                                              Human TGF-beta receptor II targeted antisense oligonucleotide #52.
                                                                                                                                                                                                              .;
o
                                                                                                                                                                                               0.8%; Score 13.8; DB 1; Length 20; 88.2%; Pred. No. 1e+03; Live 0; Mismatches 2; Indels
                                                                                                                                                                                   Sequence 20 BP; 4 A; 6 C; 7 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                           antisense oligonucleotide;
                                                                                                                                                                                                                          1631 CCAGCAGGCAGCGCTG 1647
                                                                                                                                                                                                                                       CAAGCTGGCAGCGGCTG 18
                                                                                                                                                                                                                                                                         ВЪ
                                                                                                                                                                                                                                                                                                                                                     phosphorothioate backbone; ss
                                                                                                                                                                                                                                                                                                                                                                                                          19-JUN-2002; 2002WO-US019665.
                                                                                                                                                                                                                                                                                                                                                                                                                      21-JUN-2001; 2001US-00888361.
                                                                                                                                                                                                                                                                         ADC65775 standard; DNA; 20
                                                                                                                                                                                                                                                                                                   (first entry)
                                                                                                                                                                                               Query Match
Best Local Similarity 88.2
Matches 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                   (ISIS-) ISIS PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                WO2003000656-A2
                                                                                                                                                                       Factor (PEDF)
                                                                                                                                                                                                                                                                                                                                                                   Homo sapiens
                                                                                                                                                                                                                                                                                                  18-DEC-2003
                                                                                                                                                                                                                                                                                                                                                                                             03-JAN-2003
                                                                                                                                                                                                                                                                                      ADC65775;
                                                                                                                                                                                                                                                            RESULT 1545
                                                                                                                                                                                                                                                                   ADC65775,
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New compound having a sequence targeted to a nucleic acid encoding Transforming growth factor beta-receptor II, useful for preparing a composition for treating hyperproliferative disorder e.g., lung, liver,

Wyatt JR;

Murray SF,

WPI; 2003-175279/17.

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The present invention describes isolated polynucleotides (I) encoding proteins (II) from Lolium perenne and Festuca arundinacea which are active in Ligain, fructan and tannin biosynthetic pathways. Also described: (I) an isolated oligonucleotide probe or primer comprising at least 10 contiguous residues complementary to 10 contiguous residues of (I); (2) a kt comprising the oligonucleotide probe or primer; (3) a genetic construct comprising (I); (4) a transgenic plant cell comprising the genetic construct of (3); (5) a plant or its seed, fruit or progeny comprising the transgenic plant cell of (4); (6) modulating one or more of the lignin, fructan or tannin compositions of a plant; (7) producing a plant having one or more of the lignin, fructan or tannin compositions;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New polynucleotide encoding polypeptides from Lolium perenne or Festuca arundinacea, useful for modulating the biosynthesis of lignin, fructan or tannin in a plant.
                                                           The invention comprises antisense oligonucleotides that are targeted to the nucleic acid encoding transforming growth factor beta (TGF-beta) receptor II. The antisense oligonucleotides of the invention are useful for treating: hyperproliferative disorders (e.g. breast cancer), or an autoimmune disorder (e.g. rhematoid arthritis). The present DNA sequence represents a 2'-O-methoxyethyl gapmer oligonucleotide with a phosphorothioate backbone that is targeted to human TGF-beta receptor II.
                                                                                                                                                                                                                                                                            Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Glenn M;
                                                                                                                                                                                                                                                                            ..
0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Lolium perenne; Festuca arundinacea; lignin; fructan; tannin;
biosynthetic pathway; plant; PCR primer; ss.
                                                                                                                                                                                                                                        0.8%; Score 13.8; DB 1; Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Shenk MA, Norriss MG,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Tannin biosynthesis gene related PCR primer SEQ ID NO:217.
                                                                                                                                                                                                                                                                            Indels
                                                                                                                                                                                                       Sequence 20 BP; 8 A; 4 C; 7 G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                          5
                                                                                                                                                                                                                                                            Pred. No. 1e+03;
                                                                                                                                                                                                                                                                              0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example 8; SEQ ID NO 217; 240pp; English.
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                                                                                                                                                                                                                                                                                                                1202 CCCTCTTTCCGGGCTCC 1218
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               & DEV CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                      ADC68507 standard; DNA; 20 BP.
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                                                                                                                                                                                                                                                            88.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
                                                                                                                                                                                                                                                                                15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Schedonorus arundinaceus.
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(WRIG-) WRIGHTSON SEEDS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Forster RL,
KM, Hall C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2003-441544/41.
                                                                                                                                                                                                                                               Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WO2003040306-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     lium perenne.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           18-DEC-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Synthetic.
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                                                                                                                                                                                                                                                                                                                                                                                                         RESULT 1546
                                                                                                                                                                                                                                                                                   Matches
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and (8) modifying the activity of (II) involved in a lignin, fructan or tannin biosynthetic pathway in a plant. (I) can be used for modulating the biosynthesis of lignin, fructan or tannin in a plant. The present sequence is used in the exemplification of the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (e.g. CPRA, CPRB or CYP52A1A), useful for producing dicarboxylic acids that may be utilized as industrial intermediates in manufacturing
                                                                                                                                                            Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 reductase enzymes
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The invention relates to an isolated nucleic acid selected encoding Candida tropicalis omega oxygenase complex enzymes (cytochrome P450
                                                                                                                                                              ..
                                                                                                                      Score 13.8; DB 1; Length 20;
Pred. No. 1e+03;
0; Mismatches 2; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New nucleic acid encoding cytochrome P450 and NADPH
                                                                                                                                                                                                                                                                                                                                                                                                                     Yeast CYP52A5A/B genes 5' region RT-PCR primer #1.
                                                                                     Sequence 20 BP; 6 A; 3 C; 8 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Eshoo M,
Gleeson M;
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Loper JC,
                                                                                                                                                          0;
                                                                                                                                                                                             851 TGGACAAGGACCTGAAG 867
                                                                                                                                                                                                                             TGGACATGGACCAGAAG 18
                                                                                                                                                                                                                                                                                                                     BP
                                                                                                                        0.8%;
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10-MRA-1999; 99US-0123555P
30-APR-1999; 99US-00302620.
12-OCT-2001; 2001US-00976800.
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                                                                                                                                                                                                                                                                                                                   ADC45046 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                        (first entry)
                                                                                                                                                          15; Conservative
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Tang M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                diesters and polymers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MADDURI K M.
CORNETT C A.
BRENNER A A.
TANG M.
LOPER J C.
GLEESON M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CRAFT D L.
EIRICH L D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2003-777150/73
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                                                                                                                                          Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Candida tropicalis
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  US2003049821-A1.
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                                                                                                                                                                                                                                                                                                                                                     ADC45046;
                                                                                                                         Query Match
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(GLEE/)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (ESHO/)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (TANG/)
                                                                                                                                                                                                                                                                                RESULT 1547
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                                                                                                                                                          Matches
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monooxygenase (CYP) and NADPH reductase enzymes (CPR) designated CPRA, CYP52A1A, CYP52A1A, CYP52A3B, CYP52A3B, CYP52A3B, CYP52A3B, CYP52A5B, CYP5A5B, CYP5A5B,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      diesters and polymers (e.g. as thermoplastics, plasticising agents, lubricants, hydraulic fluids, agricultural chemicals, pharmaceuticals, dyes, surfactants or adhesives). The present sequence is a quantitative competitive reverse transcription (QC-RT) PCR primer used to assay the levels of CYP, CPR or control POX mRNA in response to exogenously added
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PCR; Primer; ss; yeast; omega oxygenase complex; cytochrome P450 monooxygenase; CYP; NADPH reductase enzymes; CPR; CPRA; CYPSAB, CYPSABA; CYPSABA; CYPSABA; CYPSABA; CYPSABA; CYPSABA; CYPSABA; CYPSABA; CYPSABA; CYPSABB; CAPACABB; CAPSABB; CAPSABB;
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0
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Pred. No. 1e+03;
0; Mismatches 2; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 20 BP; 7 A; 3 C; 9 G; 1 T; 0 U; 0 Other;
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98US-0103099P.
99US-0123555P.
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88.2%;
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12-OCT-2001; 2001US-00976800.
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Best Local Similarity 88.2
Matches 15, Conservative
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CORNETT C A.
BRENNER A A.
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LOPER J C.
GLEESON M.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             substrates.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  g,
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(EIRI/)
(ESHO/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (MADD/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (BREN/)
(TANG/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              RESULT 1548
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Cornett CA;

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10-DEC-2001; 2001US-00006430
                                                                                                                   10-DEC-2001; 2001US-00006430
                                                                                                                                                     (ISIS-) ISIS PHARM INC
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              US2003113914-A1
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                                                19-JUN-2003
                                                                                                                                                                                       Graham MJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADC84236;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
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Hsu H,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1550
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         RESULT 15
ADC84236
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           dd
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       à
                                                                                                                                                                                                  condidator tropicalis omega oxygenase complex enzymes (cytochrome P450 monooxygenase (CYP) and NADPH reductase enzymes (cytochrome P450 monooxygenase (CYP) and NADPH reductase enzymes (CYR) designated CPRA, CYP52ABA, CYP52ABA, CYP52ABA, CYP52ABA, CYP52ABA, CYP52ABB, CYP5ABB, CYP5ABB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Antisense, ss, human; CD81; TAPA-1; tetraspanin; viral infection; cocaine addiction; autoimmune disorder; antiinflammatory; antibacterial; virucide; antiparasitic; inflammatory disorder; parasitic infection; bacterial infection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /mod_base= OTHER
/note= "Phosphorothioate backbone and all cytidines are
-methyl cytidines"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CPR or control POX mRNA in response to exogenously added
                                                                                   dicarboxylic acids manufacturing
                                                                       reductase enzymes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
                                                                                                                                                                                         The invention relates to an isolated nucleic acid selected encoding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     .;
0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 13.8; DB 1; Length 20;
Pred. No. 1e+03;
0; Mismatches 2; Indels
                                                                 New nucleic acid encoding cytochrome P450 and NADPH (e.g. CPRA, CPRB or CYP52A1A), useful for producing that may be utilized as industrial intermediates in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /note= "2'-methoxyethyl nucleotide"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /mod_base= OTHER
/note= "2'-methoxyethyl nucleotide"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 20 BP; 7 A; 3 C; 9 G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human CD81/TAPA-1 antisense oligonucleotide #60.
 Gleeson M;
                                                                                                                                                         Example 11; SEQ ID NO 47; 196pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /*tag= c
/mod_base= OTHER
 Loper JC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1010 AGAGGGGAGAGCTCAAG 1026
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AGAGGCAGGCTCAAG 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADC35600 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
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                                                                                                                         diesters and polymers.
 Tang M,
                                 WPI; 2003-765370/72.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity
Matches 15; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   levels of CYP,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    modified base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      modified base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              18-DEC-2003
   Brenner AA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        substrates.
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The invention relates to a compound (antisense oligonucleotide)
hybridising with the eighth nucleobase portion of an active site on a
nucleic acid molecule encoding CD81 (also known as TAPA-1, a tetraspanin)
and inhibiting the expression of CD81. Also included is a composition
comprising the antisense oligonucleotide and a carrier or a diluent. The
antisense oligonucleotide is useful for inhibiting the expression of CD81
in cells or tissues. The antisense oligonucleotide is also useful for
treating infections preferably viral, bacterial and parasitic and
diseases such as inflammatory disorders and autoimmune disorders. The
disease or condition is characterised by chemical dependency (e.g.
                                                                                         inhibiting the expression of CD81, useful for treating infections and disease associated with expression of CD81 such as inflammation disorder.
                                                          Novel compound hybridizing with nucleic acid molecule encoding CD81 and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human papillomavirus type 6 (HPV 6) detection oligonucleotide #2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             probe; human papilloma virus; HPV; detection; identification; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ΰ
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0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0.8%; Score 13.8; DB 1; Length 20;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 10. 10+03;
Pred. No. 10+03;
2; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 20 BP; 6 A; 4 C; 8 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Lee
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Pan C,
                                                                                                                                                                         Claim 3; SEQ ID NO 72; 55pp; English.
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Kao Y,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (KING-) KING CAR FOOD IND CO LTD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1700 ACTCTCTGCCTACCTGC 1716
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      88.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  17 ACTCTCTGCCTTCATGC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADC84236 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Best_Local Similarity 88.2
Matches 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Lin R, You C,
Shih C, Yeh C,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      cocaine addiction). The oligonucleotide of the
WPI; 2003-810907/76.
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                                                                                                                                   subtypes of human papilloma virus (HPV) contained in a sample. The digonucleotides of the invention are useful for simultaneously detecting and identifying subtypes of HPVs. The present DNA sequence represents an HPV detection oligonucleotide of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Detector for identifying human papilloma virus subtypes, comprises carrier having two parts carrying first and second oligonucleotides that respectively hybridize with DNA contained in first and second subtypes of
         Detector for identifying human papilloma virus subtypes, comprises carrier having two parts carrying first and second oligonucleotides that respectively hybridize with DNA contained in first and second subtypes of
                                                                                                                     The invention comprises oligonucleotides for detecting and identifying
                                                                                                                                                                                                                                                                              Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human papillomavirus type 6 (HPV 6) detection oligonucleotide #1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          probe; human papilloma virus; HPV; detection; identification; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Fan C;
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0
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                                                                                                                                                                                                                                              0.8%; Score 13.8; DB 1; Length 20;
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                                                                                                                                                                                                                                                                            2; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 20 BP; 5 A; 8 C; 1 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                               Sequence 20 BP; 6 A; 7 C; 1 G; 6 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                            Pred. No. 1e+03;
); Mismatches
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Pan C, (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 4; SEQ ID NO 465; 221pp; English.
                                                                                      Claim 4; SEQ ID NO 466; 221pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Huang H,
Kao Y,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (KING-) KING CAR FOOD IND CO LTD.
                                                                                                                                                                                                                                                                           0;
                                                                                                                                                                                                                                                                                                       1677 CCCCAACTACATCTTCC 1693
                                                                                                                                                                                                                                                                                                                                        13
                                                                                                                                                                                                                                                                                                                                                                                                                   ВР
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                                                                                                                                                                                                                                                            88.28;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human papillomavirus type 6.
                                                                                                                                                                                                                                                                                                                           ADC84235 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           You C, 1
Yeh C,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2003-432398/41.
                                                                                                                                                                                                                                                            Local Similarity
ses 15; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Lin R, Shih C,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               01-JAN-2004
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      che virus
                                                           the virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                ADC84235
                                                                                                                                                                                                                                               Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Lin C,
Hsu H,
                                                                                                                                                                                                                                                            Best Loca
Matches
                                                                                                                                                                                                                                                                                                                                                                                                     ADC84235
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The invention relates to a novel angiogenesis inhibitor comprising a compound that inhibits the activity of an amino acid sequence given in the specification. Angiogenesis-related proteins Bv8, ZAQ and ISE were utilised within the method of the invention. The molecules of the invention demonstrate cytostatic and antiinflammatory activities whilst the method may be useful for treatment and prevention of cancer, ovarian diseases, diabetic retinopathy and inflammatory disease. The current sequence is that of the angiogenesis inhibitor-related PCR primer of the
                                                                                                                                                                                                 angiogenesis inhibitor; cytostatic; antiinflammatory; cancer; ovarian disease; diabetic retinopathy; inflammatory; ZAQ; Bv8; I5E; ss; PCR; primer; RBv8-WR2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Angiogenesis inhibitors for treatment and prevention of cancer, ovarian diseases and inflammatory disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0.8%; Score 13.8; DB 1; Length 20; 88.2%; Pred. No. 1e+03; 1ive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       primer; male infertility; infertility-associated mutation;
azoospermia factor; Y-chromosome;
                                                                                                                                                                               Angiogenesis inhibitor-related PCR primer RBv8-WR2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 20 BP; 3 A; 7 C; 3 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human infertility associated primer SEQ ID 73.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example 3; SEQ ID NO 35; 308pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                       Takatsu Y;
1677 CCCCAACTACATCTTCC 1693
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              878
                                                                                                BP
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                                                                                                                                                                                                                                                                                                                                                                                14-FEB-2002; 2002JP-00027299.
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                                                                                                                                                                                                                                                                                                                                                                                                           (TAKE ) TAKEDA CHEM IND LTD
                CCGTAACTACATCTTCC
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                                                                                               standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                       Masuda Y,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2003-646310/61
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Les 15; Conserv
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                                                                                                                                                                                                                                                                   Inidentified
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                                                                                                                          ADD69057;
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Best Local Similarity 88.2 Matches 15; Conservative

Query Match

ADE28941;

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This invention describes a novel method for investigating genetic infertility or predisposition in males. The method involves selecting at least two infertility-associated mutations which are recessive or intermediate that are associated mutations which are recessive or intermediate that are associated with infertility in the heterozygous state and/or only in the homozygous state. Preferably at least one azoospermia factor is detected which may be lost by macrodaletions in intervals. Isted, present in the cystic fibrosis transmembrane conductance regulator (CTRX), Kallmann syndrome (KAL1), androgen conductance regulator (CTRX), Kallmann syndrome (KAL1), androgen conductance regulator (CTRX), Kallmann syndrome (KAL1), androgen conductancy strands, are fixed to a carrier, particularly as a microarray, then tested for hybridization with oligomucleotides from or synthesized from, a patient sample and hybridization detected. Whitifactorial analysis is by standard statistical methods, particularly the quantitative trait locus method. The method is used to diagnose inherited male infertility or predisposition to its, especially in conjunction with in vitro fertilization programs, e.g. for assessing intracycoplasmic sperm injection method. Analysis of many mutations intracycoplasmic sperm injection method. Analysis of many mutations intracycoplasmic sperm injection method. Analysis of many mutations of intracycoplasmic sperm injection method. Analysis of many mutations of intracycloides that represent known mutations (nonsense or missence, consequence analyzing genetic infertility in meles comprises of inservations, allelic variants deletions or rearrangements) in the cyclic through analyzing genetic infertility in method. Analyzing analyzing genetic analyzing genetic analyzing genetic analyzing genetic analyzing analyzing
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Investigating male genetic infertility, useful for diagnosis e.g. for assessing suitabilty for in vitro fertilization, based on multifactorial analysis of infertility-related mutations.
cystic fibrosis transmembrane conductance regulator; CTFR;
Kallmann syndrome; KAL1; androgen resistance; steroid 21-hydroxylase;
CYP21; microarray; quantitative trait locus; in vitro fertilization;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 13; SEQ ID NO 73; 110pp; German.
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                                                                                                                                                                                                       oligospermia; ss
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                                                                                                                                                                                                                                                                                                                           Homo sapiens
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Gaps . 0 Score 13.8; DB 1; Length 20; Pred. No. 1e+03; 2; Indels Sequence 20 BP; 6 A; 3 C; 7 G; 4 T; 0 U; 0 Other; 0.8%; Query Match 0.8 Best Local Similarity 88.2 Matches 15; Conservative

0

577 GTCAGCCTATCTGAGAT 593 20 GGCAGCCTATGTGAGAT

ADE28941/c ID ADE28941 standard; DNA; 20 BP. RESULT 1554

antisthmatic; antilipaemic; metabolic; diabetes; obesity, infectious; ancoexia; cancer; cardiovascular; hypertension; atherosclerosis; neurodegenerative; Alzheimer's disease; Parkinson's; epilepsy; immune; osteoarthritis; haemopoietic; inflammatory skin; asthma; dyslipidaemia; neurogenesis; cell differentiation; proliferation; haemopoiesis; wound healing; anglogenesis; gene therapy; chromosome mapping; tissue typing; human; NOV; PCR; primer; ss; RT-PCR. NOVX; antidiabetic; anorectic; cardiant; hypotensive; antiatreriosclerotic; virucide; antibacterial; fungicide; protozoacide; nootropic; neuroprotective; antipatkinsonian; anticonvulsant; osteopathic; antiatritic; antiinflammatory; dermatological; Reverse Ag2597 RT-PCR primer used to amplify human NOV RNA 12-DEC-2001; 2001US-0341346P.
17-DEC-2001; 2001US-0341477P.
17-DEC-2001; 2001US-0341540P.
27-DEC-2001; 2001US-03442592P.
31-DEC-2001; 2001US-0344903P.
17-APR-2002; 2002US-0333288P. 15-MAY-2002; 2002US-03809B1P. 17-MAY-2002; 2002US-0381495P. 28-MAY-2002; 2002US-0381534P. 28-MAY-2002; 2002US-0383744P. 29-MAY-2002; 2002US-038329P. 29-MAY-2002; 2002US-0383829P. 2002US-0401788P. 2002US-0401788P. 2002US-0406353P. 05-DEC-2001; 2001US-0336600P. 07-DEC-2001; 2001US-0338285P. 05-NOV-2002; 2002WO-US035536 31-OCT-2002; 2002US-00287971 (first entry) WO2003040330-A2. Homo sapiens. 05-NOV-2001; 07-AUG-2002; 29-JAN-2004 15-MAY-2003.

(CURA-) CURAGEN CORP.

Twomlow N; Eisen A; Mcqueeney K; Patturajan M; Shimkets RA; Alsobrook JP, Alvarez E, Anderson DW, Baron M, Boldog FL; Burgess CE, Casman SJ, Chapoval A, Dhanabal M, Edinger SR, Ellerman K, Ettenberg S, Gangolli EA, Gerlach VL, Gorman L; Grosse WM, Guo X, Hackett C, Ji W, Kekuda R, Kiramtsov NV; Lepley DW, Li L, Macdougall JR, Malyankar UM, Mazur A, Moqu Mezes PS, Miller CE, Millet I, Mishra VS, Padigaru M, Pattu Pena CEA, Peyman JA, Rastelli L, Rieger DK, Shenoy SG, Shim Smithson G, Starling G, Spytek KA, Stone DJ, Tchernev VT, T Zhong M; Peyman JA, Rast, Starling G, S Vernet CAM,

WPI; 2003-441555/41.

New isolated NOVX polypeptides and polynuclectides, useful for preventing, diagnosing or treating NOVX-associated disorders, e.g. osteoarthritis, obesity, atherosclerosis, cancer, Parkinson's disease, asthma, or infections.

Example C; SEQ ID NO 318; 447pp; English.

The invention relates to a novel isolated NOVX polypeptide. The polypeptide of the invention demonstrates, antidiabetic, anorectic, cardiant, hypotensive, antiarteriosclerotic, virucide, antibacterial, fungicide, protozoacide, nootropic, neuroprotective, antiparkinsonian, anticonvulsant, osteopathic, antiarthritic, antiinflammatcory, dermatological, antiasthmatic and antilipaemic activities. The

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polypeptides, nucleic acid molecules and antibodies may be useful for treating or diagnosing diseases including metabolic disorders such as diabetes and obesity, infectious diseases, anorexia, cencer, cardiovascular diseases including hypertension and atherosclerosis, neurodegenerative disorders such as Alzheimer's disease, Parkinson's disease and epilepsy, immune disorders, asthma and atherosclerosis, disease and epilepsy, immune disorders, asthma and dyslipidaemia. Furthermore, the nucleic acids and polypeptides may also be used to identify molecules that modulate or inhibit neurogenesis, cell differentiation and proliferation, haemopoiesis, wound healing and angiogenesis, as well as in gene therapy. Finally, the nucleic acids may be used as hybridisation probes, in chromosome mapping, tissue typing, preventive medicine and pharmacogenomics. The current sequence is that of the RT-PCR primer which was used within the exemplification of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                   invention.
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Sequence 20 BP; 10 A; 3 C; 5 G; 2 T; 0 U; 0 Other;

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                              Gaps
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 0.8%; Score 13.8; DB 1; Length 20;
88.2%; Pred. No. 1e+03;
ve 0; Mismatches 2; Indels
                            Indels
                                                     1256
             88.28;
                                                    1240 ITCAICTICCGIAICTI
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             Best_Local Similarity 88.2
Matches 15; Conservative
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Query Match
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C. tropicalis CYP52A5A/B QC-RT-PCR primer #1. BP ADE52127 standard; DNA; 20 29-JAN-2004 (first entry) ADE52127; RESULT 1555 ADE52127

Yeast; BS; PCR; primer; cytochrome P450; CYP; NADPH reductase; CPR; omega-hydroxylase complex; omega-oxidation; long chain fatty acid; QC-RT PCR; Quantitative competitive reverse transcriptase PCR.

Candida tropicalis

US2003073220-A1.

17-APR-2003

03-MAY-2002; 2002US-00138916.

98US-0083798P. 98US-0103099P. 99US-0123555P. 01-MAY-1998; 05-0CT-1998; 10-MAR-1999;

99US-00302620 12-OCT-2001; 2001US-00976800 30-APR-1999;

WILSON C R. (WILS/) CRAF/)

CRAFT D L. EIRICH L D. ESHOO M. EIRI/) ESHO/) (MADD/)

MADDURI K M. CORNETT C A. BRENNER A A. TANG M. LOPER J C. GLEESON M. (BREN/) (TANG/) (LOPE/) (GLEE/) (CORN/)

Madduri KM, Eshoo M, Gleeson M; Eirich LD, Loper JC, Craft DL, Tang M, AA, g, Brenner Wilson

Cornett CA;

WPI; 2003-625522/59.

New cytochrome P450 and NADPH oxidoreductase, i.e. CPR and CYP, genes and proteins, useful for discriminating members of a gene family by

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The invention relates to isolated nucleic acids encoding cytochrome P450 (CYP) and NADPH reductase (CPR) enzymes of the omega-hydroxylase complex of Candida tropicalis. Also included are the CYP and CPR proteins (Comprising CPRA, CPR52A1A, CYP52A2A, CYP52A2B, CYP52A3B, CYP52A5A, CYP52A3B, CYP52A5B, CYP52A5B, CYP52A3B, CYP52A5B, CYP52A5B, CYP52A3B, CYP52A5B, CYP5A5B, CYP5A5B
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quantifying the amount of target mRNA in a sample, or for omega-oxidation of long chain fatty acids.
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Pred. No. 1e+03;
0; Mismatches 2; Indels
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                                                                                                        Example 11; SEQ ID NO 47; 194pp; English.
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Best Local Similarity
Thes 15; Conserva
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detection; human papillomavirus; HPV subtype; probe; ss. BP HPV 6 detecting probe M0601. ADF44137 standard; DNA; 20 (first entry) 12-FEB-2004 ADF44137; ADF4413

RESULT 1556

28-NOV-2001; 2001JP-00362595. Human papillomavirus type 6. JP2002360271-A 17-DEC-2002

WPI; 2003-600935/57.

(KING-) KING CAR FOOD IND CO LTD

04-MAY-2001; 2001TW-00110785.

A detecting apparatus and a detecting method for identifying the subtypes of many species of human papilloma viruses at the same time and a composition for the detection.

Claim 1; SEQ ID NO 494; 166pp; Japanese.

subtypes of human papillomaviruses (HPV) contained in a sample which comprises a carrier which can load sample, a first oligonucleotide loaded on first part of the carrier and a second oligonucleotide loaded on second part of carrier, in which first and second oligonucleotides hybridise with the DNA of the first and the second HPV subtype and can identify HPV subtype contained in sample at the same time. ADF43644-This invention describes a novel detecting apparatus for identifying the

RESULT 1557

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ADF44138

Matches

555×8

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The present invention relates to a method for judging the genotype of hepatitis B viruses (HBV) in which part of the gene sequence of the PreSl region of HBV is amplified by PCR using labelled primers and the amplified product is hybridized with HBV type A, B, C, D, B, F and G gene specific probes and the label in the PCR product is detected.
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                                                                                                                                                                                                                                                                                                                            Classification of genotype of hepatitis B viruses and primers and probes for the \operatorname{method}.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match

0.8%; Score 13.8; DB 1; Length 20;
Best Local Similarity 88.2%; Pred. No. 1e+03;
Matches 15; Conservative 0; Mismatches 2; Indels
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                                SEQ ID 9.
                                                                                                                                                                                                                                                                                                                                                                                    Claim 3; Page 3; 13pp; Japanese.
                                                                                                                                                                                                                                                                 (GENO-) GENOME SCI KENKYUSHO KK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1058 CAATCCCAACAAGACA 1074
                                Hepatitis B virus PreS1 probe,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    17
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12-FEB-2004 (first entry)
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MADDURI K M.
CORNETT C A.
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BIRICH L D.
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                                                                                                Hepatitis B virus
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                                                                   PreS1; HBV;
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(CRAF/)
(EIRI/)
(ESHO/)
(MADD/)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     This invention describes a novel detecting apparatus for identifying the subtypes of human papillomaviruses (HPV) contained in a sample which comprises a carrier which can load sample, a first oligonucleotide loaded on first part of the carrier and a second oligonucleotide loaded on second part of carrier, in which first and second oligonucleotides subtype and can identify HPV subtype contained in sample at the same time. ADP$43644-ADP$44289 represent oligonucleotide probes used in the method of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         subtypes
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                                                                                                                       Gaps
   ADF44289 represent oligonucleotide probes used in the method of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       A detecting apparatus and a detecting method for identifying the of many species of human papilloma viruses at the same time and a composition for the detection.
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                                                                                                                       Indels
                                                                                                                                                                                                                                                                                                                                                                                                       detection; human papillomavirus; HPV subtype; probe; ss.
                                                                                    DB 1; Length
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                                                     BP; 5 A; 8 C; 1 G; 6 T; 0 U; 0 Other;
                                                                                      Score 13.8; DB 1
Pred. No. 1e+03;
                                                                                                                         0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 1; SEQ ID NO 495; 166pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (KING-) KING CAR FOOD IND CO LTD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1677 CCCCAACTACATCTTCC 1693
                                                                                                                                                        1693
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADF70749 standard; DNA; 20 BP.
                                                                                                                                                                                       20
                                                                                                                                                                                                                                                                        BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          28-NOV-2001; 2001JP-00362595
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          04-MAY-2001; 2001TW-00110785
                                                                                        0.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                           ģ
                                                                                                                                                                                                                                                                                                                                                                         HPV 6 detecting probe M0602
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1677 CCCCAACTACATCTTCC
                                                                                                                                                                                                                                                                        ADF44138 standard; DNA; 20
                                                                                                                                                                                       4 ccgraacracarcricc
                                                                                                                                                                                                                                                                                                                                        (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                           Human papillomavirus type
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
                                                                                                                         15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2003-600935/57
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local Similarity
Matches 15; Conserv
                                                                                        Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            JP2002360271-A.
                                                         Sequence 20
                                                                                                                                                                                                                                                                                                                                          12-FEB-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           17-DEC-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 invention.
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Gaps

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ADF70749

ADE70749 ID ADE7 XX AC ADE7 XX

RESULT 1558

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GLEE/)
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                                                                                                                                                The invention describes an isolated nucleic acid encoding a CYP52A2B protein comprising the fully defined sequence of 522 amino acids, as given in the specification, and comprising a coding region defined by nucleotides 1072-2640 of a fully defined sequence of 3755 base pairs, as given in the specification. The nucleic acids encoding the cytochrome P450 and NADH reductase enzymes of Candida tropicalis are useful for increasing the production of dicarboxylic acid for oxidising organic substrates such as fatty acids. This sequence represents a quantative competition reverse transcriptase PCR (QC-RT-PCR) primer for quantating the level of Candida tropicalis CYP52A5 RNA in a sample.
                                                                                                         93
                                                                                           New nucleic acids encoding a CYP52A2B protein useful for increasing th
production of dicarboxylic acid for oxidizing organic substrates such
                                                Cornett CA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                           ss; primer; QC-RT-PCR; CPRA; CPRB; CYP52A1A; CYP52A2A; CYP52A2B; CYP52A3A; CYP52A3B; CYP52A5B; CYP52A5B; CYP52A5B; CYP52D4A; gene family; quantitative competitive reverse transcription.
                                                                                                                                                                                                                                                                                                Gaps
                                                                                                                                                                                                                                                                                                 ·,
                                                                                                                                                                                                                                                                           0.8%; Score 13.8; DB 1; Length 20; 88.2%; Pred. No. 1e+03; ive 0; Mismatches 2; Indels
                                               Madduri KM,
                                                                                                                                                                                                                                                          Sequence 20 BP; 7 A; 3 C; 9 G; 1 T; 0 U; 0 Other;
                                               LD, Eshoo M,
                                                                                                                                Example 10; SEQ ID NO 47; 188pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                        C. tropicalis QC-RT-PCR primer #11.
                                              Eirich LD
Loper JC,
                                                                                                                                                                                                                                                                                                                 1010 AGAGGGGAGAGCTCAAG 1026
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    98US-0103099P.
99US-0123555P.
99US-00302620.
                                                                                                                                                                                                                                                                                                                                  AGAGGGCAGGCTCAAG 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           98US-0083798P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       03-MAY-2002; 2002US-00139218
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 12-OCT-2001; 2001US-00976800
                                                                                                                                                                                                                                                                                                                                                                                 ADF11874 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                        (first entry)
                                                                                                                                                                                                                                                                                                Conservative
                                              Craft DL,
                                                       Tang M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MADDURI K M.
CORNETT C A.
BRENNER A A.
BRENNER A A.
                                                                          WPI; 2003-810780/76.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CRAFT D L.
EIRICH L D.
ESHOO M.
         TANG M.
LOPER J C.
GLEESON M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WILSON C R.
                                                                                                                                                                                                                                                                                     Local Similarity
tes 15; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Candida tropicalis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   US2003153060-A1.
                                                                                                              fatty acids.
                                                        Brenner AA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           01-MAY-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     05-OCT-1998;
10-MAR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      30-APR-1999;
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(BREN/) (TANG/) (LOPE/)
                                                                                                                                                                                                                                                                            Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                     ADF11874;
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(BREN/)
                          GLEE/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (EIRI/)
(ESHO/)
                                              Wilson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (MILS/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (CRAF/)
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                                                                                                                                                                                                                                                                                              Matches
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                                                                                                                                                                                                                                                                                   Cornett CA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ss; primer; QC-RT-PCR; CPRA; CPRB; CYP52A1A; CYP52A2A; CYP52A2B; CYP52A3A; CYP52A3B; CYP52A4A; GYP52A4A; GYP5CA4B; CYP5CA4B; CYP5CA4B; CYP5CA4B; GYP5CA4B; GYP5CAAB; G
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CYP52A5A, CYP52A5B, CYP52ABA, CYP52A2B, CYP52A3B, CYP52A5B, CYP5A5B, CYP5
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1e+03;
                                                                                                                                                                                                                                                                                   Madduri KM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 20 BP; 7 A; 3 C; 9 G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                            , Eshoo M,
Gleeson M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0.8%; Score 13.8; E88.2%; Pred. No. 1e+0
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                                                                                                                                                                                                                                                                            Eirich LD,
Loper JC, G
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1010 AGAGGGGAGAGCTCAAG 1026
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05-OCT-1998; 98US-0103099P.
10-MAR-1999; 99US-0123555P.
301-APR-1999; 99US-00305620.
12-OCT-2001; 2001US-00976800.
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                                                                                                                                                                                                                                                                            Craft DL,
Tang M,
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ESHOO M.
MADDURI K M.
CORNETT C A.
BRENNER A A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2003-897719/82.
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ses 15; Conserv
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(TANG/) TANG M.
(LOPE/) LOPER J C.
(GLEE/) GLEESON M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CRAFT D L.
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LOPER J C.
GLEESON M.
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                                                                                                                                                                                                                                                                                                                                                   Brenner AA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         12-FEB-2004
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                                                                                                                                                                                                                                                                     Wilson CR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADF11756;
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(EIRI/)
(ESHO/)
(MADD/)
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(TANG/)
(LOPE/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (MILS/)
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is complementary to a second variant nucleotide at the position, thus to require missincorporation of the first nucleotide at the position, thus to reduce missincorporation of the first nucleotide on a template comprising molecular events for identifying individuals with mutations indicative of cancer. They are particularly useful in detecting a rare mutation in a heterogeneous biological sample (e.g. sporadic mutation in a heterogeneous biological sample), detecting rare genotypes in genotyping reactions (e.g. viral genotyping reactions), or detecting mutant or viral sequences in pooled samples (e.g. detecting polymorphisms or inherited sequence variations in pooled patient samples). This polymucleotide sequence represents a primer used as part of the primer extension assay
nucleotide that is complementary to a first variant nucleotide suspected
                                                                                                                                                                                                                                                                                                                      the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RESULT 1563
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Matches
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    $$$$$$$$$$$$$$$$$$$$
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                                                                                                                                                                                                                                                                    The invention relates to a new isolated nucleic acid which encodes a CPRA, CPRS_ASA, CYPS_ASA, CASA, CAS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The invention relates to a novel method for detecting a variant in a primer extension assay, useful for analysing molecular events for identifying mutations indicative of cancer, by contacting a target nucleic acid primer complementary to a region of the target nucleic acid primer complementary to a region of the target nucleic acid the target nucleic acid primer complementary to a region of the target nucleic acid primer complementary to a region of the target nucleic acid primer complementary to a region of the target nucleic acid contacting a nucleic acid, and extending the primer in the presence of a first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Detecting a variant in a primer extension assay, useful for analyzing molecular events for identifying mutations indicative of cancer, by contacting a target nucleic acid primer complementary to a region of the target nucleic acid.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Variant detecting primer extension assay extension primer, SEQ ID No 24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    variant detection; primer extension assay; mutation; cancer;
heterogeneous; sporadic mutation; genotyping; pooled sample; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
         ď.
                                                                                                                         CPRB, CYP52A1A, CYP52A2A, CYP52A2B, CYP52A3A, CYP52A3B, CYP52A5B, CYP52A5B, CYP52A8B or CYP52D4A gene, useful for
         Cornett
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 13.8; DB 1; Length 20; Pred. No. 1e+03; 0; Mismatches 2; Indels
       Madduri KM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Seguence 20 BP; 7 A; 3 C; 9 G; 1 T; 0 U; 0 Other;
       Eirich LD, Eshoo M, oper JC, Gleeson M;
                                                                                                                                                                                                                          Example 11; SEQ ID NO 47; 196pp; English
                                                                                                                                                                                discriminating members of a gene family.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 3; SEQ ID NO 24; 54pp; English.
                                Loper JC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1010 AGAGGGGGGGGGCTCAAG 1026
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match 0.8%;
Best Local Similarity 88.2%;
Matches 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2 AGAGGGCAGGGCTCAAG
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                                  Tang M,
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                                                                               WPI; 2003-897579/82.
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                                  Brenner AA,
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            Wilson CR,
                                                                                                                              CPRA,
                                                                                                                                                    CYP52A5A,
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                                                                                                                                                                                                                                                                                                                                                                                                                    primer.
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                                                                                                                       Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  human; single nucleotide polymorphism; microarray; side effect; ss;
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                                                          DB 1; Length 20;
                                                                                                                       Indels
Sequence 20 BP; 7 A; 5 C; 3 G; 5 T; 0 U; 0 Other;
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                                                      Score 13.8; DB 1
Pred. No. 1e+03;
                                                                                                                    Mismatches
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                                                             0.8%;
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                                                                                                                             Conservative
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                                                                                          Local Similarity
es 15; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            primer; PCR
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                                                          Query Match
Best Local S
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first active agent comprising an oligonuclectide antisense to the intitation codon, coding region, 5' or 3' end genomic flanking regions, 5' and 3' intron-exon junctions, or regions within 2-10 nuclectides of junctions of genes encoding a polypeptide associated with lung and/or nacal airway dysfunction and a second active agent comprising an antiinflammatory steroid and ubiquinone. A composition of the invention has antiinflammatory, antiallergic, antistshmatic, hypotensive, immunosuppressive, and cytostatic activity. The composition may have a use in antisense gene therapy. The composition is useful for treating or preventing a respiratory, lung or malignant disease or condition, also for enhancing the prophylactic or therapeutic respiratory effect of an antisinal manarory steroid in a subject, for reducing levels of adenosine contiinflammatory steroid in a subject, for reducing levels of adenosine continuing surfactant in a subject, st issue, or treating bronchoconstriction, lung allergies, or a respiratory disease or condition, lung allergies, or a respiratory disease or condition. Note: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Pharmaceutical composition for treating ailments associated with impaired
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human, antisense; lung dysfunction, nasal airway dysfunction; antiinflammatory; antiallergic; antiinflammatic; hypotensive; immunosuppressive; cyfostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy;
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                                                                             DB 1; Length 20;
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le+63;
le+63;
le+63;
                                      Sequence 20 BP; 3 A; 1 C; 11 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     lung inflammation; respiratory disease; ds.
                                                                                                                0; Mismatches
                                                                           Score 13.8;
Pred. No. 1e
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Tang L, Shahabuddin S;
detection method of the invention,
                                                                                                                                                                                                                                                                                                                                                                                                              Human oligonucleotide sequence.
                                                                                                                                                       227 AGAGTGGTGGTGGC 243
                                                                                                                                                                                           4 AGATTGGTGGAGGTGGC 20
                                                                                                                                                                                                                                                                                              ABZ86270 standard; DNA; 20 BP
                                                                           0.8%;
88.2%;
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                                                                                                                15; Conservative
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                                                                                             Local Similarity
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The invention relates to a novel pharmaceutical composition, which has a first active agent comprising an oligonucleotide antisense to the initiation coodon, coding region, 5' or 3' end genomic flanking regions, 5' and 3' intron-exon junctions, or regions within 2-10 nucleotides of junctions of genes encoding a polypeptide associated with lung and/or nasal airway dysfunction and a second active agent comprising an antiinflammatory steroid and ubjquinone. A composition of the invention has antiinflammatory, antiallergic, antiasthmatic, hypotensive, immunosuppressive, and cytostatic activity. The composition may have a use in antisense gene therapy. The composition is useful for treating or preventing a respiratory, lung or malignant disease or condition, also for enhancing the prophylactic or therapeutic respiratory effect of an preventing sensitivity to adenosine. reducing or depleting levels of, or reducing sensitivity to adenosine, reducing levels of adenosine receptor, producing pronchodilation, increasing levels of ubiquinone or ung surfactent in a subject's tissue, or treating bronchoconstriction, lung inflammation, lung allergies, or a respiratory disease or condition. Note: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or
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                                                                                                                          Gaps
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                                                                              0.8%; Score 13.8; DB 1; Length 20; 88.2%; Pred. No. 1e+03; tive 0; Mismatches 2; Indels
                                        Sequence 20 BP; 4 A; 4 C; 5 G; 7 T; 0 U; 0 Other;
at ftp.wipo.int/pub/published_pct_sequences
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Tang L, Shahabuddin S;
                                                                                                                                                                                                                                                                                                                                                                                                                             Human oligonucleotide sequence.
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                                                                                                                                                              58 TGACTGCTGAAACCCAG
                                                                                                                                                                                                      19 TGACTGCTGAAATACAG
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Query Match
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Miller S,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human; antisense; lung dysfunction; nasal airway dysfunction; antiinflammatory steroid; ubiquinone; antiinflammatory; antiallergic; antiasthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy;
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                                                                                                                                         Score 13.8; DB 1; Length 20;
Pred. No. 1e+03;
0; Mismatches 2; Indels
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                                                                                 Sequence 20 BP; 6 A; 6 C; 2 G; 6 T; 0 U; 0 Other;
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              at ftp.wipo.int/pub/published pct sequences
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i, Shahabuddin
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                                                                                                                                                    0.8%;
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                                                                                                                                                                                                                                                                                                 343 TTGAAGATGGGGTCTGA
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                                                                                                                                            Query Match 0.8 Best Local Similarity 88.2 Matches 15; Conservative
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Tang L,
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Miller S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ABZ97631;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              lung inflammation, lung allergies, or a respiratory disease or condition. Note: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO
                                                                                                                                                                                                                                                                                                                                                                                                                           Human, antisense; lung dysfunction, nasal airway dysfunction, antiinflammatory steroid; ubiquinone; antiinflammatory; antiallergic; antiinflammatic; hypotensive; ortostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy; lung inflammation; respiratory disease; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The invention relates to a novel pharmaceutical composition, which has a first active agent comprising an oligonucleotide antisense to the initiation codon, coding region, 5 or 3 end genomic flanking regions, 5 and 3' intron-exon junctions, or regions within 2-10 nucleotides of junctions of genes encoding a polypeptide associated with lung and/or antiallary dysfunction and a second active agent comprising an antinflammatory steroid and ubiquinone. A composition of the invention has antinflammatory antiallergic, antiasthmatic, hypotensive, immunosuppressive, and cytostatic activity. The composition may have a use in antisense gene therapy. The composition is useful for treading or preventing a respiratory, lung or malignant disease or condition, also for enhancing the prophylactic or therapeutic respiratory effect of an
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               antinflammatory sferoid in a subject, for reducing or depleting levels of, or reducing sensitivity to adenosine, reducing levels of adenosine receptor, producing bronchodilation, increasing levels of ubiquinone or lung surfactant in a subject's tissue, or treating bronchoconstriction, lung inflammation, lung allergies, or a respiratory disease or condition
                                                                              Gaps
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0
                                        Length 20;
                                                                              Indels
Sequence 20 BP; 6 A; 4 C; 4 G; 6 T; 0 U; 0 Other;
                                                                            7
                                        Score 13.8; DB 1;
Pred. No. 1e+03;
                                                                              0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Li Y, Sandrasagra A,
Tang L, Shahabuddin
                                                                                                                         1444 ATGAAACATCCATTCTT 1460
                                                                                                                                                                                                                                                                                                                                                                                         Human oligonucleotide sequence
                                                                                                                                                             19
                                                                                                                                                                                                                                                                      BP.
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                                        0.8%;
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                                                                                                                                                         3 ATGAAGCATCCATACTT
                                                                                                                                                                                                                                                                      ABZ91330 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                    (first entry)
                                                                                15; Conservative
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                                                             Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Homo sapiens.
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                                        Query Match
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                                                                                   Matches
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Tue Nov

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Gaps

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DB 1; Length 20; :+03; s 2; Indels

Sequence 20 BP; 2 A; 3 C; 9 G; 6 T; 0 U; 0 Other;

at ftp.wipo.int/pub/published_pct_sequences

SXS

Query Match 0.8%; Score 13.8; DB 1 Best Local Similarity 88.2%; Pred. No. 1e+03; Matches 15; Conservative 0; Mismatches

548 ACAAGCCCCTCAGCCGC 564

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7

18 ACAAGGCCCTCAACCGC

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first active agent comprising an oligomortal contribution within the first active agent comprising an oligomortal contribution codon, coding region, 5' or 3' end genomic flanking regions, 5' and 3' intron-exon junctions, or regions within 2-10 nucleotides of junctions of genes encoding a polypeptide associated with lung and/or nasal airway dysfunction and a second active agent comprising an entile flammatory steroid and ubiquinone. A composition of the invention has antiinflammatory, antiallergic, antiasthmatic, hypotensive, immunosuppressive, and cytostatic activity. The composition may have a use antiinflammatory und or malignant disease or condition, also for enhancing the prophylactic or therapeutic respiratory effect of an antiinflammatory steroid in a subject, for reducing or depleting levels of on the inflammatory sensitivity to adenosine, reducing levels of adenosine receptor, producing bronchodilation, increasing levels of adenosine receptor, producing bronchodilation, increasing bronchoconstriction, lung surfactant in a subject's tissue, or treating bronchoconstriction, lung inflammation, lung allergies, or a respiratory disease or condition.

Specification, but was obtained in electronic format directly from WIPO
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human, antisense; lung dysfunction, nasal airway dysfunction, antiinflammatory; steroid, ubiquinone; antiinflammatory; antiallergic; antisathmatic; hypotensive; immunosuppressive, cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy;
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                                                                                                                          Gaps
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                                                                                    DB 1; Length 20;
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                                                                                        1;
1e+03;
2;
                                          Sequence 20 BP; 3 A; 6 C; 8 G; 3 T; 0 U; 0 Other;
at ftp.wipo.int/pub/published_pct_sequences
                                                                                                                        0; Mismatches
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                                                                                  0.8%; Score 13.8;
88.2%; Pred. No. 1e
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Li Y, Sandrasagra A, Ka
Tang L, Shahabuddin S;
                                                                                                                                                                1269 TGAGGAGACGTGGCCAG 1285
                                                                                                                                                                                                                                                                                                                                                                                                                                Human oligonucleotide sequence.
                                                                                                                                                                                                        4 TGAGGACACGTGGCCTG 20
                                                                                                                                                                                                                                                                                                          ABZ93366 standard; DNA; 20 BP
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                                                                                                                          15; Conservative
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                                                                                                       Local Similarity
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Miller S,
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                                                                                                                                                                                                                                                                                                                                                  ABZ93366;
                                                                                Query Match
                                                                                                                          Matches
    SXS
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preventing a respiratory, lung or malignant disease or condition, also for enhancing the prophylactic or therapeutic respiratory effect of an anti-inflammatory steroid in a subject, for reducing or depleting levels of, or reducing sensitivity to adenosine, reducing levels of adenosine receptor, producing bronchodilation, increasing levels of ubiquinone or lung surfactant in a subject's tissue, or treating bronchoconstriction, lung inflammation, lung allergies, or a respiratory disease or condition. Note: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Pharmaceutical composition for treating ailments associated with impaired
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The invention relates to a novel pharmaceutical composition, which has a first active agent comprising an oligonucleotide antisense to the initiation codon, coding region, 5' or 3' end genomic flanking regions, 5' and 3' intron-exon junctions, or regions within 2-10 nucleotides of junctions of genes encoding a polypeptide associated with lung and/or nasal airway dysfunction and a second active agent comprising an antiinflammatory steroid and ubiquinone. A composition of the invention has antiinflammatory, antiallergic, antiasthmatic, hypotensive, immunosuppressive, and cytostatic activity. The composition may have a use in antisense gene therapy. The composition is useful for treating or
                                                                                                                                                                                           Human, antisense, lung dysfunction, nasal airway dysfunction, antiinflammatory steroid, ubiquinone; antiinflammatory, antiallergic, antiasthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy, antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy; lung inflammation; respiratory disease; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Pabalan J, Aguilar D;
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S;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 15; SEQ ID NO 992; 872pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Li Y, Sandrasagra A,
Tang L, Shahabuddin
                                                                                                                                                  Human oligonucleotide sequence.
                    BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (EPIG-) EPIGENESIS PHARM INC.
                  ABZ85750 standard; DNA; 20
                                                                                                          (first entry)
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Miller S,
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ABZ85750,
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88.2%;

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ABZ80343
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human; retinal pigmented epithelium derived neurotrophic factor; PEDF; retinal disease; retinal tumour; retinoblastoma; retinal detachment; neuronal-retinal tumour; macular degeneration; retinitis pigmentosa; diabetic retinopathy; inherited and age-related pathology; tumour; ocular disease; nerve injury; serine protease related disorder; cytostatic; ophthalmological; antiinflammatory; antidiabetic; PCR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Treating retinal disease such as retinal tumors, retinitis pigmentosa,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      involves
                                                                                                                                                                              Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                macular degeneration and diabetic retinopathy, in a subject, in administering Pigment Epithelium Derived Factor to the subject.
                                                                                                                                                                           ;
                                                                                                               0.8%; Score 13.8; DB 1; Length 20; 88.2%; Pred. No. 1e+03;
                                                                                                                                                                           Indels
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                                                                                                                                                                        7
                                                            BP; 6 A; 7 C; 2 G; 5 T; 0 U; 0 Other;
at ftp.wipo.int/pub/published_pct_sequences
                                                                                                                                                                           Mismatches
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0
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92US-00952796.
94US-00279979.
95US-00377710.
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                                                                                                                                                                                                                                                                                          17 AGATGGGCCTGTATGAG 1
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                                                                                                                                                                                                                                    212 AGATAGGCCTGGATGAG
                                                                                                                                                                                                                                                                                                                                                                                                                                       ABS57272 standard; DNA; 20
                                                                                                                                                                           Conservative
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                                                                                                            Query Match
Best Local Similarity
Matches 15; Conserv
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24-SEP-1992;
25-JUL-1994;
25-JAN-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Homo sapiens.
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                                                            Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   30-JAN-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   primer; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ABS57272;
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DB 1; Length 20;

0.8%; Score 13.8;

Query Match

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The present invention describes a method (M) for generating a substantially homogeneous population of undifferentiated cells (UC) from substantially homogeneous population of undifferentiated cells (UC) from a biological sample (BS), which comprises subjecting BS or its sub-sample to tissue-disruption to provide a mixed population (MP) comprising UC. corsequentially with SD, subjecting the cell population obtained to a cell size-discrimination step, Also described: (1) a cell-surface marker-discrimination step, Also described: (1) a cell-surface marker-discrimination step, Also described: (1) a comprising a population of undifferentiated cells (I) comprising a prowth factor identified using a homogeneous population of neural stem cells (NSCs) generated by (M); and (3) a composition (III) comprising a growth factor identified using a homogeneous population of neural stem cells (NSCs) generated by (M). (1) can have nootropic, neuroprotective and antiparkinsonian activities, and can be used in gene therapy. (M) is useful for generated by (M). (1) can have nootropic, neuroprotective and antiparkinsonian activities, and can be used in gene therapy. (M) is useful for the replacement of neural or non-neural tissue in an animal. (II) is useful in cell replacement therapy in an organ such as the brain or treating a CNS disorder such as Albitaimer's disease, treating a CNS disorder such as Albitaimer's disease, parkinson's disease, acute brain injury and CNS dysfunction. (I) is useful for the replacement such as Albitaimer's disease, parkinson of tissue. ABZ80278 to ABZ80363 represent for markers defining cell populations
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Generating substantially homogeneous population of undifferentiated cells from sample, by disrupting tissue sample, discriminating cells in population based on size and performing cell-surface marker-
                                                                                                                                                                                                                                                                                                                                                                                                     neuroprotective, antiparkinsonian, gene therapy, nervous system, central nervous system, CNS, Alzheimer's disease, Parkinson's disease, acute brain injury, CNS dysfunction, tissue regeneration, tissue repair,
                     Gaps
                                                                                                                                                                                                                                                                                                                                                                                   Purification; neural stem cell; NSC; undifferentiated; nootropic;
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0
                        Indels
                     2;
Pred. No. 1e+03;
                                                                                                                                                                                                                                                                                                                                           Mouse Emx1 antisense PCR primer SEQ ID NO:66.
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                        Mismatches
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                                                                 1647
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                                                                                                                                                                                                                  BP.
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                                                                                                           2 CAAGCTGGCAGCGGCTG
                                                                                                                                                                                                                  ABZ80343 standard; DNA; 20
                                                                    1631 CCAGCAGGCAGCGGCTG
                                                                                                                                                                                                                                                                                                    (first entry)
    Best Local Similarity 88.2
Matches 15, Conservative
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                                                                                                                                                                                                                                                                                                    28-MAY-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PCR primer;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Synthetic.
                                                                                                                                                                                                                                                            ABZ80343;
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(MONI/) MONIA B P.
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                                                                                                                                                                                                                                                                                                                                                                                    25-JAN-2002;
                                                                                                                                                                                                                                                                                                                    Unidentified
                                                                                                                                                                                                                                                                                                                                                                                                                                                               28-AUG-1998;
18-FEB-2000;
                                                                                                                                                                                                                     05-SEP-2003
                                                                                                                                                                                                                                                                                                                                                                                                         31-MAY-1994;
                                                                                                                                                                                                                                                                                                                                                              13-FEB-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                     31-MAY-1995;
26-NOV-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                    06-JUL-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                         07-JUL-1997
                                                                                                                                                                                                 ACD42154;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Monia BP;
                                                       Query Match
                                                                   Best Local
                                                                                                                                                      RESULT 1573
                                                                            Matches
                                                                                                                                                                ACD42154/
  88833
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                                                                                                                     셤
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New antisense chimeric oligonucleotide, useful for modulating the expression of human Interleukin 12 p40 subunit, in treating or preventing disease states in humans and animals, and as research reagents and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The invention relates to an antisense compound 20-50 nucleobases in length targeted to a start codon region, coding region, a stop codon region or a 3'-untranslated region of a nucleic acid molecule encoding human Interleukin 12 p40 subunit. The compound specifically hybridises with one of the regions and inhibits the expression of human Interleukin 12 p40 subunit. The new compound is useful for inhibiting the expression of human Interleukin 12 p40 subunit in cells or tissues and comprises contacting the cells or tissues in vitro with the compound, so that expression of the human Interleukin 12 p40 subunit is inhibited. The antisense compound may also be used as research reagents and diagnostics, and as treatment or prevention of disease states, e.g. to prevent or
                                                                                                                                                                                                         Human interleukin 12 p40 subunit antisense oligonucleotide ISIS #139149.
                                                                                                                                                                                                                                                                                                                           /note= "All cytosines are 5-methylcytidines and the nucleotides are linked via a phosphorothioate backbone"
                                            Gaps
                                                                                                                                                                                                                              Human, 88; antisense; interleukin 12 p40 subunit; antibacterial;
antiinflammatory; cytostatic; infection; inflammation; tumour.
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                                                                                                                                                                                                                                                                                                                                                                      /mod_base= OTHER
/note= "2'-methoxyethyl (2'-MOE) nucleotides"
16. .20
                                                                                                                                                                                                                                                                                                                                                                                                                           /note= "2'-methoxyethyl (2'-MOE) nucleotides"
                     Score 13.8; DB 1; Length 20; Pred. No. 1e+03;
                                           Indels
 Sequence 20 BP; 1 A; 10 C; 1 G; 8 T; 0 U; 0 Other;
                                           2;
                                           Mismatches
                                                                                                                                                                                                                                                                                    Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 15; Col 45; 42pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                  base= OTHER
                                                                                                                                                                                                                                                                                                                   OTHER
                                           ;
0
                                                                935
                                                                                   4 rrccrcrrccAGCTTCT 20
                                                                                                                                           BP
                     0.8%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           07-MAY-2001; 2001US-00851062
                                                               919 TICCIGITCCAGCIGCT
                                                                                                                                        ABX33976 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                   /mod base=
                                                                                                                                                                                                                                                                                                      *tag= a
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                                           Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Baker BF, Freier SM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2003-074100/07.
                              Local Similarity
es 15; Conserv
                                                                                                                                                                                                                                                                                   Key
modified_base
                                                                                                                                                                                                                                                                                                                                                                                            modified base
                                                                                                                                                                                                                                                                                                                                                  modified base
                                                                                                                                                                                                                                                              sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                  US6448081-B1
                                                                                                                                                                                   10-FEB-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     diagnostics
                                                                                                                                                              ABX33976;
                    Query Match
                                                                                                                     RESULT 1572
                                                                                                                                                                                                                                                             Homo
                                           Matches
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The invention relates to an oligonucleotide 8-50 nucleotides in length which is targeted to mRNA encoding human c-raf, a-raf or b-raf (raf is a protein kinase playing a regulatory role in signal transduction, regulating cell proliferation and has been implicated in lung carcinoma), and which is capable of inhibiting raf expression. Also included is a composition comprising the oligonucleotide and a pharmaceutically acceptable carrier. The antisense oligonucleotide is useful for inhibiting the expression of human raf in human cells or tissues, by contacting the human cells or tissues with the oligo. The oligo. is also is useful for treating or preventing a disease or condition associated with the expression of raf by administering it in combination with a chemotherapeutic agent to a human or cells of the human, where the chemotherapeutic agent to a human or cells of the human, where the chemotherapeutic agent to a human or cells of the human, where the chemotherapeutic agent to a human cannot an or cells of the numan, where the chemotherapeutic of raf is abnormal expression, and the condition is a human new angiogenesis or neovascularisation (preferrably ocular angiogenesis or neovascularisation). The oligo. is also useful for inhibiting
                                                                                                                                                                                                                                                                                                      ö
delay infection, inflammation or tumour formation, in animals and humans. The present sequence is an antisense oligonucleotide of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Novel antisense oligonucleotide which is targeted to mRNA encoding human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Antisense; c-raf; a-raf; b-raf; protein kinase; cancer; ss; signal transduction; cell proliferation; lung carcinoma; cytostatic; antisense gene therapy; chemotherapeutic agent; angiogenesis; hyperproliferative condition; neovascularisation; ocular angiogenesis.
                                                                                                                                                                                                                                                                                                          Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         raf and which is capable of inhibiting raf expression, useful for treating or preventing hyperproliferative conditions such as cancer.
                                                                                                                                                                                                                                                                                                      0;
                                                                                                                                                                                                                                                                                                      2; Indels
                                                                                                                                                                                                         0.8%; Score 13.8; DB 1; Length:
8.2%; Pred. No. 1e+03;
ve 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human raf-associated antisense oligonucleotide #16.
                                                                                                                               Sequence 20 BP; 4 A; 9 C; 3 G; 4 T; 0 U; 0 Other,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Disclosure; Page 32; 42pp; English.
                                                                                                                                                                                                                                                                                                                                                                                     480 ACTACCAGCTGACATCC 496
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95WO-US007111.
96US-00756966.
97US-00888982.
98WO-US013961.
98US-01014314.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           BP
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                                                                                                                                                                                                                                                     88.2%;
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ACD42154 standard; DNA; 20
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                                                                                                                                                                                                                                                                                                 15; Conservative
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                                                                                                                                                                                                                                                              Similarity
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hyperproliferation of cells. The oligos. are also useful as tools, for example for detecting and determining the role of raf expression in various cell functions and physiological processes and conditions and for diagnosing conditions associated with raf expression and for research purposes. The present sequence is an antisense oligonucleotide included in the sequence listing but not mentioned elsewhere in the specification
                                                                                                                                                                                                                                                                                                                                               nootropic, antiulcer, muscular, immunosuppressive, gynaecological, antipsoriatic, endocrine, ophthalmological; osteopathic; antiparkinsonian, anticonvolsant; tranquiliser, analgesic, nephrotropic; antiinfertility; antilipaemic; cardiomyopathy; athersselerosis; diabetes; cell signal processing; metabolic pathway, asthma, allergy; emphysema; autoimmune; graft-versus-host; arthritis; cancer; stroke; haemophilia;
                                                                                                                                                                                                                                                                                                    NOVX; cytostatic; antiarteriosclerotic; antidiabetic; antiasthmatic; antiallergic; antiinflammatory; respiratory; antiarthritic; dermatological; antibacterial; cerebroprotective; vasotropic; cardiant; haemostatic; hypotensive; hepatotropic; neuroprotective; anorectic;
                                                                                                                                                                                                                                                                                                                                                                                                               obesity, Alzheimer's; pain; chromosome mapping; tissue typing; human;
RT-PCR; PCR; primer; ss; NOV8.
                                                                                                                     Gaps
                                                                                                                    ;
                                                                                                                                                                                                                                                                                 RT-PCR primer used to analyse human NOV8 expression - SEQ 171,
                                                                                             0.8%; Score 13.8; DB 1; Length 20;
8.2%; Pred. No. 1e+03;
ve 0; Mismatches 2; Indel8
                                                                           Sequence 20 BP; 6 A; 10 C; 0 G; 4 T; 0 U; 0 Other;
                                                                                                                                         1152 TGACATGTGGGTGTGG 1168
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20010S-0274122P
20010S-0274826P
20010S-0279840P
20010S-0282981P
20010S-028364P
20010S-0393666P
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2000US-0258171P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2001US-0311754P
                                                                                                          88.28;
                                                                                                                                                        17 TGAGATGTGTGTGTGG
                                                                                                                                                                                                                    ADJ83180 standard; DNA; 20
                                                                                                                                                                                                                                                            (first entry)
                                                                                                                    15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SHIMKETS R A. GROSSE W M. SZEKERES E S. VERNET C A M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SPYTEK K A.
ZERHUSEN B D.
PATTURAJAN M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ALSOBROOK J P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ICHERNEV V T.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       BURGESS C E.
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                                                                                                          Local Similarity
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11-APR-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens.
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08-MAR-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           22-MAR-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          13-APR-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     31-JUL-2001;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          17-AUG-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 21-DEC-2000;
22-DEC-2000;
                                                                                                                                                                                                                                                             06-MAY-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        11-SEP-2003
                                                                                                                                                                                                                                         ADJ83180;
                                                                                                Query Match
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(SZEK/)
(VERN/)
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currentiation, designate and controlled and control
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New NOVX polypeptides and nucleic acid molecules, useful for diagnosing, preventing or treating NOVX-associated polypeptide disorder, e.g. cardiomyopathy, atherosclerosis, diabetes, cancer, Parkinson's disease or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The invention relates to a novel isolated NOVX polypeptide. The polypeptide demonstrates cytostatic, antiatrareziosotlerotic, antidiabetic, antiasthmatic, antiallergic, antiallargic, antiallargic, antiallargic, antiallargic, antiplamatory, respiratory, antiarthritic, dermatological, antibacterial, cerebroprotective,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Grosse WM;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Smithson G;
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18.2%; Pred. No. 1e+03;
ve 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Tchernev VT, Liu X, Spytek KA, Zerhusen Lepley DM, Burgess CE, Shinmkers RA, Grobernet CAM, Lii LGSaman SJ, Boldog FL, Fernandes ER, Rieger DK, Edinger SR, Gun!
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 20 BP; 10 A; 3 C; 5 G; 2 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example 2; SEQ ID NO 171; 263pp; English.
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es 15; Conservative
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В
                                                                                                                                                                                  FERNANDES E R.
RIEGER D K.
LI L.
CASMAN S J.
BOLDOG F L.
GORMAN L.
GANGOLLI E A.
                                                                                                                                                                                                                                                                EDINGER S.R.
GUNTHER E.
                                                                                                                                                                                                                                                                                                                                      MILLET I.
SCIORE P.
ELLERMAN K.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SMITHSON G.
                                                                                                                                                                                                                                                                                                                                                                                                                                                   MACDOUGALL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Patturajan M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Alsobrook JP.
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Gangolli EA,
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(ELLE/)
(MACD/)
   (LILL/)
                                                                                                                                                                                      (FERN/)
(RIEG/)
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                                                                                                                                                       GANG/)
                                                                                                                                                                                                                                                                (EDIN/)
                                                                                                            GORM/)
                                                                                                                                                                                                                                                                                                       (COUNTY)
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                                                                             BOLD/
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADM83655,
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The invention describes a method of diagnosing a cellular proliferative disorder of breast tissue in a subject comprising determining the state of methylation of one or more nucleic acid isolated from the subject, where the state of methylation of one or more nucleic acids is compared with the state of methylation of one or more nucleic acids from a subject of the state of methylation of one or more nucleic acids from a subject of the amount of the state of methylation of one or more nucleic acids from a subject of the activities of the methylation state of cyclin 14-3-3 sigma Opd islands in commal mammary epithelium, breast cancer cell lines and in primary
predisposition; reverse transcriptase PCR; RT-PCR; primer; CpG island; ss; cyclin 14-3-3 sigma; human.
                                                                                                                                                                                                                                                                                                                                                                                                           Diagnosing a cellular proliferative disorder of breast tissue in subject comprises determining the state of methylation of one or nucleic acid isolated from the subject.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 20 BP; 3 A; 9 C; 3 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 12; SEQ ID NO 42; 59pp; English.
                                                                                                                                                                                                                                                                                                                                                  Dooley WC,
                                                                                                                                                                                                   26-JAN-2001; 2001US-00771357.
                                                                                                                                                                   28-JAN-2002; 2002US-00059579
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       88.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                                                                                                                                                                                  EVRON E.
DOOLEY W C.
SACCHI N.
DAVIDSON N.
                                                                                                                                                                                                                                                                                                                                              Sukumar S, Evron E,
                                                                                                                                                                                                                                                                                                                                                                              WPI; 2003-851722/79.
                                                                                                                                                                                                                                                                                                                (FACK/) FACKLER M J.
                                                                                                                                                                                                                                   SUKUMAR S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Local Similarity
es 15; Conserv
                                                                                                   US2003138783-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          mammary tumours.
                                                                        Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
                                                                                                                                                                                                                                 (SUKU/)
                                                                                                                                                                                                                                                                                (SACC/)
(DAVI/)
                                                                                                                                                                                                                                                   EVRO/)
                                                                                                                                                                                                                                                                   (DOOI/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Matches
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a more

Sacchi N, Davidson N, Fackler MJ;

Gaps ·. DB 1; Length 20; 2; Indels le+03; 0.8%; Score 13.8; DI 38.2%; Pred. No. 1e+0: ive 0; Mismatches

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843 TGAGTACCTGGACAAGG 859
                                    ABD30662 standard; DNA; 20 BP
          N
          TGAGTACCGGGAGAAGG
                                               ABD30662;
                          RESULT 1576
                                ABD30662
                                     ठे
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Human; antisense; bronchoconstriction; allergy; hyposecretion; pain; respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antiinflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis;

Human IL5-R derived oligonucleotide SEQ ID 12873.

(first entry)

29-JUL-2004

RESULT 1577 ABD29596/c

This invention describes a novel composition (a) a first active agent, comprising oligonuclectides, effective for alleviating comprising oligonuclectides, effective for alleviating by the proposition of proposecretion, when administered to a mammal. The reducing adenosine sensitivity, levels of adenosine (A) or (A) receptors, surfactant depletion or hyposecretion, when administered to a mammal. The oligonuclectides are derived from a gene encoding or regulating dysfunction or cancer and can be anti-sense to the corresponding mRNA. The invention also describes a kit, that comprises: (a) a delivery device, in separate containers, (b) the oligonuclectides, (c) instructions for adding a carrier and for use of the kit. The composition of the invention has antiallergic, antiinflammatory, antiasthmatic, analgesic, hypotensive, immunosuppressive and cytostatic activity, is a beta-adrenergic agonist. The composition is useful for preventing or treating a respiratory, lung or malignant disease. The administered composition comprises oligo and is administered to reduce the production or availability, or to increase the degradation of the target mRNA or to reduce the amount of target polypeptide present in the lungs. The reduce the amount of target polypeptide present in the lungs. The culmonary obstruction, and/or surfactant hypoproduction are associated with a disease or condition such as pulmonary vasoconstriction, and/or surfactant hypoproduction and sortion of inflammation, allergies and/or surfactant hypoproduction and sortion of distress syndrome, pain, cystic fibrosis, allergic rhinitis, pulmonary infections, bronchitis or cancer. The reduced adenosine content of the anti-sense oligos corresponding to thymidines present in the target RNA serves to prevent the breakdown of the oligonucleotides into products that free adenosine into the system of eg., lung, brain, heart, kidney, etc, tissue environment and thereby, to respiratory distress syndrome; allergic rhinitis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis; pulmonary transplantation rejection; ss; primer. beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; Pharmaceutical composition for treating asthma, has antisense oligonuclectide containing less percentage of adenosine, targeted to nucleic acids associated with lung airway or lung dysfunction, and Pabalan J, Aguilar D; Sequence 20 BP; 6 A; 4 C; 4 G; 6 T; 0 U; 0 Other; Claim 15; SEQ ID NO 12873; 763pp; English. Katz E, prevent any unwanted effects due to it Li Y, Sandrasagra A, Ka Tang L, Shahabuddin S; 23-APR-2002; 2002WO-US013143. 24-APR-2001; 2001US-0286036P. (EPIG-) EPIGENESIS PHARM INC bronchodilating agent. WPI; 2003-093058/08. WO200285309-A2 31-OCT-2002. Nyce JW, Miller S,

0 0.8%; Score 13.8; DB 1; Length 20; 18.2%; Pred. No. 1e+03; Indels .; ? 0; Mismatches 1444 ATGAAACATCCATTCTT 1460 19 88.2%; 3 Argaagcarccaracrr 15; Conservative Local Similarity Query Match Matches ð

ö

Gaps

ABD29596 standard; DNA; 20

ABD29596

(first entry) 29-JUL-2004 H86812-derived oligonucleotide SEQ ID 8608.

analgesic; hypotensive, immunosuppressive; cytostatif; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory distress syndrome; allergic rininitis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis; pulmonary transplantation rejection; ss; primer. Human; antisense; bronchoconstriction; allergy; hyposecretion; pain; respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antiinflammatory; antiasthmatic;

Homo sapiens.

31-OCT-2002

23-APR-2002; 2002WO-US013143.

24-APR-2001; 2001US-0286036P

(EPIG-) EPIGENESIS PHARM INC

Aguilar D; Pabalan J, Katz E, s, Sandrasagra A, Li Y, Sar Tang L, Nyce JW,] Miller S,

WPI; 2003-093058/08.

t oligonucleotide containing less percentage of adenosine, targeted touches acids associated with lung airway or lung dysfunction, and bronchodilating agent. Pharmaceutical composition for treating asthma, has antisense

Claim 15; SEQ ID NO 8608; 763pp; English.

This invention describes a movel composition, allergies and bronchoconstriction, respiratory tract inflammation, allergies and bronchoconstriction, respiratory tract inflammation, allergies and bronchoconstriction, respiratory tract inflammation, allergies and creducing adenosine sensitivity, levels of adenosine (A) or (A) receptors, surfactant depletion or hyposecretion, when administered to a mammal. The oligonucleotides are derived from a gene encoding or regulating comparison of a target polypeptide associated with lung airway or lung dysfunction or cancer and can be anti-sense to the corresponding mRNA. The invention also describes a kit, that comprises: (a) a delivery device, in separate containers, (b) the oligonucleotides, (c) instructions for adding a carrier and for use of the kit. The composition of the invention has antiallergic, antiinflammatory, antiasthmatic, analgesic, hypotensive, immunosuppressive and cytostatic activity, is a beta-adrenergic agonist. The composition is useful for preventing or treating a respiratory, lung or malignant disease. The administered composition comprises oligo and is administered to reduce the production or availability, or to increase the degradation of the target mRNA or to reduce the amount of target polypeptide present in the target mRNA or to reduce the amount of target polypeptide present in the target mRNA or confidence to inflammation, allergies, asthma, impeded respiration, respiratory distress syndrome, pain, cystic fibrosis, allergic shrintiss pulmonary therefore demosine content of the anti-sense oligos corresponding to the minimation, emphysema, chronic obstructive pulmonary disease, pulmonary infections, bronchitis or cancer. The reduced adenosine content of the anti-sense oligos corresponding to the minimal or the substant man of the anti-sense or prevent the breakdown of the minimal or the system of the anti-sense or condition products that free adenosine but of the anti-sense or condition products that free adenosine of the anti-sense or condition invention describes a novel composition (a) a first active agent prevent any unwanted effects due to it

Sequence 20 BP; 2 A; 3 C; 9 G; 6 T; 0 U; 0 Other;

0; Gaps 0 Length 20; Indels , , DB 1; Score 13.8; DB 1 Pred. No. 1e+03; Mismatches 0, 564 0.8%; 548 ACAAGCCCCTCAGCCGC Conservative Best Local Similarity 15; Query Match Matches à

18 ACAAGGCCCTCAACCGC

g

ABD21980,

ABD21980 standard; DNA; 20

ABD21980;

BP.

(first entry) 29-JUL-2004 Human stanniocalcin-derived oligo SEQ ID 992.

Human, antisense; bronchoconstriction, allergy; hyposecretion; pain; respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antiinflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cycostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory distress syndrome; allergic rhinitis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis; pulmonary transplantation rejection; ss; primer.

Homo sapiens.

WO200285309-A2

31-OCT-2002.

23-APR-2002; 2002WO-US013143.

24-APR-2001; 2001US-0286036P.

(EPIG-) EPIGENESIS PHARM INC.

Aguilar D; Pabalan J, Katz E, ŝ Li Y, Sandrasagra A, Tang L, Shahabuddin Nyce JW, Miller S,

WPI; 2003-093058/08.

10 oligonucleotide containing less percentage of adenosine, targeted nucleic acids associated with lung airway or lung dysfunction, and bronchodilating agent. Pharmaceutical composition for treating asthma, has antisense

Claim 15; SEQ ID NO 992; 763pp; English.

comprising oligonucleotides, effective for alleviating active agent, comprising oligonucleotides, effective for alleviating bronchoconstriction, respiratory tract inflammation, allergies and reducing adenosine sensitivity, levels of adenosine (A) or (A) receptors, surfactant depletion or hyposecretion, when administered to a mammal. The oligonucleotides are derived from a gene encoding or regulating expression of a target polypeptide associated with lung airway or lung dysfunction or cancer and can be anti-sense to the corresponding mRNA. The invention also describes a kit, that comprises: (a) a delivery device, in separate containers, (b) the oligonucleotides, (c) instructions for adding a carrier and for use of the kit. The composition of the invention humaniallergic, antiinflammatory, antiasthmatic, and the contractions for adding a carrier and for use of the kit. The composition of the invention human antiallergic, antiinflammatory, antiasthmatic, and the contractions for adding a carrier and for use of the kit. The composition of the invention human antiallergic, antiinflammatory, antiasthmatic, and the contractions for adding a carrier and for use of the kit. The composition of the invention human antiallergic and for use of the first and for use of the first and for use of the invention human antiallergic and for use of the first and for use analgesic, hypotensive, immunosuppressive and cytostatic activity, is a beta-adrenergic agonist. The composition is useful for preventing or treating a respiratory, lung or malignant disease. The administered composition comprises oligo and is administered to reduce the production or availability, or to increase the degradation of the target mRNA or to pulmonary obstruction, and/or bronchoconstriction and/or lung inflammation, allergies and/or surfactant hypoproduction are associated This invention describes a novel composition (a) a first active agent,

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Gaps

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Length 20; 2; Indels

DB 1;

inflammation, allergies, asthma, impeded respiration, respiratory distress syndrome, pain, cystic fibrosis, allergic rhinitis, pulmonary hypertension, emphysema, chronic obstructive pulmonary disease, pulmonary transplantation rejection, pulmonary infections, bronchitis or cancer. The reduced adenosine content of the anti-sense oligos corresponding to thymidines present in the target RNA serves to prevent the breakdown of the oligonuclectides into products that free adenosine into the system e.g., lung, brain, heart, kidney, etc, tissue environment and thereby, to prevent any unwanted effects due to it Human; antisense; bronchoconstriction; allergy; hyposecretion; pain; respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antiinflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory distress syndrome; allergic rhinitis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis; pulmonary transplantation rejection; ss; primer. Pharmaceutical composition for treating asthma, has antisense oligonuclectide containing less percentage of adenosine, targeted trucleic acids associated with lung airway or lung dysfunction, and disease or condition such as pulmonary vasoconstriction, Sequence 20 BP; 6 A; 7 C; 2 G; 5 T; 0 U; 0 Other; Score 13.8; DB 1 Pred. No. 1e+03; 0; Mismatches Human cathepsin C-derived oligo SEQ ID 1512. Katz E, i Y, Sandrasagra A, Ka Tang L, Shahabuddin S; ; 212 AGATAGGCCTGGATGAG 228 ABD22500 standard; DNA; 20 BP 0.8%; AGATGGGCCTGTATGAG 1 23-APR-2002; 2002WO-US013143. 24-APR-2001; 2001US-0286036P (EPIG-) EPIGENESIS PHARM INC 29-JUL-2004 (first entry) 15; Conservative WPI; 2003-093058/08. Local Similarity Li Y, WO200285309-A2. Homo sapiens. 31-0CT-2002 Nyce JW, L Miller S, 17 ABD22500; Query Match with a **datches** ð ద

The invention also describes a kit, that comprises: (a) a delivery device, in separate containers, (b) the oligonucleotides, (c) instructions for adding a carrier and for use of the kit. The composition of the invention has antiallergic, authinflammatory, antiasthmatic, analgesic, hypotensive, immunosuppressive and cytostatic activity, is a beta-adrenergic agonist. The composition is useful for preventing or reating a respiratory, lung or maijanant disease. The administered composition comprises oligo and is administered to reduce the production or availability, or to increase the degradation of the target mRNA or to reduce the amount of target polypeptide present in the lungs. The cultumation, allergies and/or bronchoconstriction and/or lung confilmmation, allergies and/or surfactant hypoproduction are associated with a disease or condition such as pulmonary vasoconstriction, with a disease or condition such as pulmonary vasoconstriction, continued in capture of the respiratory disease, pulmonary contension, emphysema, chronic obstructive pulmonary disease, pulmonary charaplantation rejection, pulmonary infections, bronchitis or cancer. The reduced adenosine content of the anti-sense oligos corresponding to the indignucleotides into products that free adenosine into the system of the oligonucleotides into products that free adenosine into the system of e.g., lung, brain, heart, kindiney, etc., tissue environment and thereby, to .; 0 Gaps ä . 0 Aguilar Pharmaceutical composition for treating asthma, has antisense 1; Length 20; 2; Indels Pabalan J, Sequence 20 BP; 4 A; 4 C; 5 G; 7 T; 0 U; 0 Other; Query Match
0.8%; Score 13.8; DB 1
Best Local Similarity 88.2%; Pred. No. 1e+03;
Matches 15; Conservative 0; Mismatches AA504431-derived oligonucleotide SEQ ID 6572. Katz E, prevent any unwanted effects due to it Li Y, Sandrasagra A, Ka Tang L, Shahabuddin S; 74 ВЪ 23-APR-2002; 2002WO-US013143. 24-APR-2001; 2001US-0286036P. (EPIG-) EPIGENESIS PHARM INC. 19 reacrecreaaaracae 58 TGACTGCTGAAACCCAG ABD27560 standard; DNA; 20 (first entry) WPI; 2003-093058/08. WO200285309-A2. 29-JUL-2004 31-OCT-2002. Miller S, ABD27560; Nyce JW, RESULT 1580 ठ

Human; antisense, bronchoconstriction, allergy, hyposecretion, pain; respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic, antiinflammatory; antiasthematic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory distracts syndrome; allergic rhinitis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis; pulmonary transplantation rejection; ss; primer. oligonuclectide containing less percentage of adenosine, targeted to nucleic acids associated with lung airway or lung dysfunction, and

tΩ

Pabalan J, Aguilar D;

This invention describes a novel composition (a) a first active agent, comprising oligonucleotides, effective for alleviating bronchoconstriction, respiratory tract inflammation, allergies and reducing adenosine sensitivity, levels of adenosine (A) or (A) receptors, surfactant depletion or hyposecretion, when administered to a mammal. The oligonucleotides are derived from a gene encoding or resplating expression of a target polypeptide associated with lung airway or lung dysfunction or cancer and can be anti-sense to the corresponding mRNA.

Claim 15; SEQ ID NO 1512; 763pp; English.

bronchodilating agent.

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ABD25640 standard; DNA; 20 BP
                                                                                                (first entry)
                                                                                                                             WO200285309-A2
                                                                                                                         Homo sapiens
                                                                                                                                31-OCT-2002.
                                                                                                 29-JUL-2004
                                                                                             ABD25640;
                                                                                      RESULT 1581
                                                                                       ABD25640,
                                                                                           a
                                                                             à
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This interaction describes a novel comparation, alleriating bronchoconstriction, respiratory tract inflammation, alleriating bronchoconstriction, respiratory tract inflammation, alleriating bronchoconstriction, respiratory tract inflammation, allerias and to a marmal. The cattering adenosine sensitivity, levels of adenosine (A) or (A) receptors, surfactant depletion or hyposecretion, when administered to a marmal. The coligonucleotides are derived from a gene encoding or regulating expression of a target polypeptide associated with lung airway or lung dysfunction or cancer and can be anti-sense to the corresponding mRNA. The invention also describes a kit, that comprises: (a) a delivery device, in separate containers, (b) the oligonucleotides, (c) instructions for adding a carrier and for use of the kit. The composition of the invention has antiallergic, antiinflammatory, antiathmatic, is a composition canding a respiratory, lung or malignant disease. The administered composition comprises oligo and is administered to reduce the production or availability, or to increase the degradation of the target mRNA or to reduce the amount of target polypeptide present in the lungs. The composition, allergies and/or bronchoconstriction and/or lung confinements associated inflammation, allergies and/or bronchoconstriction are associated inflammation, allergies and/or surfactant hypoproduction are associated inflammation, allergies and/or surfactant hypoproduction are associated inflammation. inflammation, allergies, asthma, impeded respiration, respiratory distress syndrome, pain, cystic fibrosis, allergic rhinitis, pulmonary hypertension, emphysema, chronic obstructive pulmonary disease, pulmonary transplantation rejection, pulmonary infections, bronchitis or cancer. The reduced adenosine content of the anti-sense oligos corresponding to thymidines present in the target RNA serves to prevent the breakdown of the oligonucleotides into products that free adenosine into the system c.y., lung, brain, heart, kidney, etc, tissue environment and thereby, to prevent any unwanted effects due to it invention describes a novel composition (a) a first active agent, Gaps ., 0.8%; Score 13.8; DB 1; Length 20; 2; Indels Sequence 20 BP; 3 A; 6 C; 8 G; 3 T; 0 U; 0 Other; Pred. No. 1e+03; 0; Mismatches Claim 15; SEQ ID NO 6572; 763pp; English. 88.2%; bronchodilating agent.

1269 TGAGGAGGTGGCCAG 1285 Query Match 0.8 Best Local Similarity 88.2 Matches 15; Conservative

20 readeacacerdecerd AI024215-derived oligonucleotide SEQ ID 4652.

beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction, respiratory distress syndrome; allergic rhinitis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis; pulmonary transplantation rejection; ss; primer. analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; Human, antisense, bronchoconstriction, allergy, hyposecretion, pain, respiratory tract inflammation, adenosine sensitivity, lung, cancer, surfactant depletion, antiallergic, antiinflammatory, antiasthmatic,

23-APR-2002; 2002WO-US013143.

This invention describes a novel composition (a) a first active agent, comprising oligonucleotides, effective for alleviating comprising oligonucleotides, effective for alleviating bronchoconstriction, respiratory tract inflammation, allergies and reducing adenosine sensitivity, levels of adenosine (A) or (A) receptors, configuration or hyposecretion, when administered to a mammal. The oligonucleotides are derived from a gene encoding or regulating expression of a target polypeptide associated with lung airway or lung dystunction or cancer and can be anti-sense to the corresponding mRNA. The invention also describes a kit, that comprises: (a) a delivery device, in separate containers, (b) the oligonucleotides, (c) instructions for adding a carrier and for use of the kit. The composition of the invention has antiallergic, antisflammatory, antiathmatic, analyses; hypotensive, immunosuppressive and cytostatic activity, is a beta-adrenergic agonist. The composition is useful for preventing or beta-adrenergic agonist. The composition is useful for preventing or tracting a respiratory, lung or mallspant disease. The administered composition comprises oligo and is administered to reduce the production or availability, or to increase the degradation of the target mRNA or to reduce the amount of target polypeptide present in the lungs. The pulmonary obstruction, and/or bronchoconstriction and/or lung inflammation, allergies and/or surfactant hypoproduction are associated with a disease or condition such as pulmonary vasoconstruction, inflammation, allergies, asthma, impeded respiration, respiratory distress syndrome, pain, cystic fibrosis, allergic rhinitis, pulmonary hypertension, emphysema, chronic obstructive pulmonary disease, pulmonary e.g., lung, brain, heart, kidney, etc, tissue environment and thereby, to prevent any unwanted effects due to it transplantation rejection, pulmonary infections, bronchitis or cancer. The reduced adenosine content of the anti-sense oligos corresponding to thymidines present in the target RNA serves to prevent the breakdown of the oligonucleotides into products that free adenosine into the system Pharmaceutical composition for treating asthma, has antisense oligonucleotide containing less percentage of adenosine, targeted to nucleic acids associated with lung airway or lung dysfunction, and Aguilar D; 0.8%; Score 13.8; DB 1; Length 20; Pabalan J, Sequence 20 BP; 6 A; 6 C; 2 G; 6 T; 0 U; 0 Other; Katz E, Claim 15; SEQ ID NO 4652; 763pp; English. ŝ Li Y, Sandrasagra A, Tang L, Shahabuddin 24-APR-2001; 2001US-0286036P. (EPIG-) EPIGENESIS PHARM INC. bronchodilating agent. WPI; 2003-093058/08. Miller S, Query Match

Gaps ö 2; Indels Pred. No. 1e+03; .. 343 TTGAAGATGGGGTCTGA 359 88.2%; 15; Conservative Similarity Local Matches à

ó

ADO84907 standard; DNA; 20 20 RESULT 1582 ADO84907, g

BP.

Human BRCA 1 and BRCA 2 mutation-related oligonucleotide probe SeqID12.

(first entry)

29-JUL-2004

ADO84907;

mutation; BRCA 1; BRCA 2; diagnosis; breast cancer; ovarian cancer; detection; predisposition; susceptibility; cancer; probe; ss; human

Homo sapiens

KR2003017894-A.

WO2003031594-A2 Homo sapiens. 04-MAR-2003. 12-AUG-2004 206 ADP75344; Query Match Cho DY, RESULT 1583 Matches ADP75344 셤 HANGE TO THE TOTAL 8

ó Mutations of BRCA 1 and BRCA 2 useful for diagnosis of breast and ovarian cancers, and detection method of predisposition and susceptibility of the This invention relates to novel mutations of BRCA 1 and BRCA 2 genes useful for the diagnosis of breast and ovarian cancers. In addition, the invention also relates to a detection method of predisposition and susceptibility of the cancers using the same. The present sequence is that of an oligonucleotide probe which may be used for detection of the Gaps Yoon GS ; 0 0.8%; Score 13.8; DB 1; Length 20; 88.2%; Pred. No. 1e+03; ive 0; Mismatches 2; Indels Song GH, Sequence 20 BP; 3 A; 6 C; 4 G; 7 T; 0 U; 0 Other; BG, o human BRCA mutations of the invention. Cho HM, Kang CS, Kim JG, Claim 3; SEQ ID NO 12; 1pp; Korean. 25-AUG-2001; 2001KR-00051510 25-AUG-2001; 2001KR-00051510 Conservative cancers using the same. (MYDN-) MYDNA CO LTD. WPI; 2003-501054/47. Local Similarity les 15; Conserv

CTGAGCAGATAGGCCTG 222 19 crdágaagaragecerg 3

Human endophilin 2 gene exon F reverse sequencing primer. ADP75344 standard; DNA; 20 BP (first entry)

single nucleotide polymorphism; a disintegrin and metalloprotease with thrombospondin typel motif 2; asthma; atopy; obesity; inflammatory bowel disease; respiratory disorder. ıman; ss; primer; ADAM19; Endophilin 1; Endophilin 2; NRG2; ADAMTS2; disintegrin and metalloprotease; neuroregulin 2; SNP;

11-OCT-2002; 2002WO-US032700.

11-OCT-2001; 2001US-0328424P.

(GENO-) GENOME THERAPEUTICS CORP.

Little RD,

Keith T, Allen K;

WPI; 2003-381712/36.

New isolated nucleic acid or alternate splice variant, useful for diagnosing and treating a disintegrin and metalloprotease (ADAM) or

The invention relates to an isolated nucleic acid or alternate splice variant comprising a nucleotide sequence containing at least one of the studies of them, or complements of them. The genes are ADAM19 (a disintegrin and metalloproclasse of them, or complements of them. The genes are ADAM19 (a disintegrin and metalloproclasse of them, or complements of them. The genes are ADAM19 (a disintegrin and metalloproclasse of them thrombospondin typel motif 2, also known as gene 873), endophilin 2, also known as gene 873, endophilin 3, also known as a carrier, and at least one component to detect the bybridisation of the variant or the binding of the antibody to the interactor gene amino acid sequence), a kit contecting an ADAM or interactor gene amino acid sequence (component to detect the binding of the antibody to the interactor gene amino acid sequence (component to detect the bybridisation and ADAM or interactor gene amino acid sequence (component to detect the bybr identifying an ADAM corrector gene ligand and an isolated nucleic acid variant of Gene 803, 845, 847, 874 or 962. The nucleic acid or alternate splice variants, methods, kits and antibody/antibody fragment are useful for diagnosing and treating an ADAM or interactor generassociated disorder, e.g. asthma, atopy, obesity or inflammatory bowel The present sequence is a primer used to sequence the regions interactor gene-associated disorder, e.g. asthma, atopy, obesity or surrounding polymorphisms in the above genes. Claim 2; Page 127; 338pp; English inflammatory bowel disease.

Sequence 20 BP; 4 A; 8 C; 2 G; 6 T; 0 U; 0 Other;

Gaps .. 0.8%; Score 13.8; DB 1; Length 20; 88.2%; Pred. No. 1e+03; ve 0; Mismatches 2; Indels 88.28; 15; Conservative Local Similarity Query Match Matches

; 0

1681 AACTACATCTTCCCTGC 1697 2 AATGACATCTTCCCTGC

ò

(first entry) 29-JAN-2004 ADE64291; KW KW KE KA KK

Del Mastro RG;

Dupuis J,

Van Eerdewegh P,

BP

ADE64291 standard; DNA; 20

1584

ADE64291

C. tropicalis CYP52A5A/B QC-RT-PCR primer #1.

Yeast; ss; PCR; primer; NADPH reductase; CPR; cytochrome P450; CYP; omega-hydroxylase; dicarboxylic acid; QC-RT PCR; Quantitative competitive reverse transcriptase PCR.

10-APR-2003

(WILS/) (CRAF/)

ESHO/) EIRI/) MADD/)

(CORN/) (BREN/)

TANG/)

(LOPE/) (GLEE/)

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The invention relates to a compound targeted to a nucleic acid molecule encoding a beta-site amyloid precursor protein (APP) -cleaving enzyme. The antisense oligonucleotides and compounds are useful for inhibiting the expression of beta-site amyloid precursor protein (APP) -cleaving enzyme, modulating amyloid deposition in neurons, altering the expression of a splice variant of beta-site APP-cleaving enzyme, and for treating diseases or conditions associated with expression of beta-site APP-cleaving enzyme e.g. neurodegeneration or Alzhaimer's disease. The antisense compounds are also useful as research reagents and kits, or in diagnostic, therapeutic and prophylaxis applications, e.g. to prevent or delay infection, inflammation or tumour formation. The present sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ss; human; beta-site amyloid precursor protein; APP-cleaving enzyme; amyloid deposition; neurodegeneration; Alzheimer's disease; infection; inflammation; tumour; antisense.
                                           ss; human; beta-site amyloid precursor protein; APP-cleaving enzyme; amyloid deposition; neurodegeneration; Alzheimer's disease; infection; inflammation; tumour.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New antisense compound targeted to a nucleic acid molecule encoding a beta-site amyloid precursor protein (APP)-cleaving enzyme, useful for treating diseases associated with beta-site APP-cleaving enzyme, e.g.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human APP-cleaving enzyme antisense oligonucleotide ISIS 223841.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0.8%; Score 13.8; DB 1; Length 20; 88.2%; Pred. No. 1e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            represents a human APP-cleaving enzyme target region.
Human APP-cleaving enzyme target region ISIS 140495.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 20 BP; 4 A; 8 C; 4 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example 15; SEQ ID NO 122; 58pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          883
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              BP
                                                                                                                                                                                                                                                                                                                     31-MAY-2002; 2002US-00159942.
                                                                                                                                                                                                                                                                                                                                                                      31-MAY-2002; 2002US-00159942
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          873 CCTGGATGACTGTGGGA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADG86683 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           17 cerecarcacrereaca
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                             (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2004-051909/05
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Local Similarity
ses 15; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   neurodegeneration.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             US2003224512-Al.
                                                                                                                                                                                                                  US2003224512-A1
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                                                                                                                                                                 Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    11-MAR-2004
                                                                                                                                                                                                                                                                    04-DEC-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADG86683;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Dobie KW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             RESULT 1586
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Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADG86683
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The invention relates to an isolated CPRA, CYPS2AJA, CYPS2AJA, CYPS2AJA, CYPS2AJA, CYPS2AJA, CYPSZAJA, CALAGO, COMPISION COMPISION COMPISION CONTRIBUTION CON
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   .
0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Novel isolated CPRA, CPRB, CYP52A1A, CYP52A2A, CYP52A2B, CYP52A3A, CYP52A3B, CYP52A5B, CYP52A5B, CYP52A6B or CYP52D4A protein, useful for increasing production of dicarboxylic acid in cells.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Cornett CA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ;
0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Madduri KM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 20 BP; 7 A; 3 C; 9 G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LD, Eshoo M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example 11; SEQ ID NO 47; 195pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Eirich LD
Loper JC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1010 AGAGGGGAGAGCTCAAG 1026
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           18
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                                                                                                                                                                                                                                                   98US-0083798P.
98US-0103099P.
99US-0123555P.
99US-00302620.
                                                                                                                                                                                                                                                                                                                                                           12-OCT-2001; 2001US-00976800
                                                                                                                                                                                                  03-MAY-2002; 2002US-00138905
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADG86739 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AGAGGGCAGGGCTCAAG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Craft DL,
Tang M,
                                                                                                                                                                                                                                                                                                                                                                                                                                        CRAFT D L.
EIRICH L D.
ESHOO M.
MADDURI K M.
CORNETT C A.
BRENNER A A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2004-020205/02.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                  WILSON C R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TANG M.
LOPER J C.
GLEESON M.
                                              Candida tropicalis.
                                                                                               US2003068800-A1
                                                                                                                                                                                                                                                                                 05-OCT-1998;
10-MAR-1999;
                                                                                                                                                                                                                                                         01-MAY-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Wilson CR,
Brenner AA,
                                                                                                                                                                                                                                                                                                                                     30-APR-1999
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molecule The

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Gaps

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2; Indels

04-DEC-2003

(first entry)

11-MAR-2004

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Matches

δ 셤 1585

ADG86739, RESULT

10017621-381.rng

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The invention relates to a compound targeted to a nucleic acid molecule encoding a beta-site amyloid precursor protein (APP)-cleaving enzyme. The antisense oligonucleotides and compounds are useful for inhibiting the expression of beta-site amyloid precursor protein (APP)-cleaving enzyme, modulating amyloid deposition in neurons, altering the expression of a splice variant of beta-site APP-cleaving enzyme, and for treating classes or conditions associated with expression of beta-site APP-cleaving enzyme e.g. neurodegeneration or Alzheimer's disease. The artisense compounds are also useful as research reagents and kits, or in diagnostic, therapeutic and prophylaxis applications, e.g. to prevent or delay infection, inflammation or tumour formation. The present sequence represents a human APP-cleaving enzyme antisense oligonucleotide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0;
                                                                                                                                                                   New antisense compound targeted to a nucleic acid molecule encoding a beta-site amyloid precursor protein (APP)-cleaving enzyme, useful for treating diseases associated with beta-site APP-cleaving enzyme, e.g.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Detecting a conserved structures in an RNA sequence by generating an offspring group from the parent group and selecting at least one group from the parent and offspring groups with the highest fitness.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                detection; conserved structure; RNA structural element; fitness; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 13.8; DB 1; Length 20;
Pred. No. 1e+03;
0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Porto VW,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Ferritin related oligonucleotide Bin#3 structure 7.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 20 BP; 4 A; 4 C; 8 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Fogel GB,
                                                                                                                                                                                                                                                Example 15; SEQ ID NO 66; 58pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Griffey RH,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 873 CCTGGATGACTGTGGGA 889
                 31-MAY-2002; 2002US-00159942.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 20
                                               31-MAY-2002; 2002US-00159942
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             10-JUN-2003; 2003WO-US018573.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          10-JUN-2002; 2002US-0387342P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADH27256 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  15; Conservative
                                                                            (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (ISIS-) ISIS PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Ecker DJ,
                                                                                                                                         WPI; 2004-051909/05
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2004-062371/06.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Local Similarity
                                                                                                                                                                                                                      neurodegeneration
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WO2003104478-A2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sampath R,
                                                                                                           Dobie KW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADH27256;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RESULT 1587
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Matches
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The present invention describes a method for detecting a conserved structure in an RNA sequence. The method comprises: (a) placing 2 structures from structures generated for 2 RNA sequences from 2 organisms into a parent group; (b) generating an offspring group from the parent group; (c) determining fitness of the parent and offspring groups; (d) comparing the fitness of the parent and offspring groups; and (e) selecting at least one group from the parent and offspring groups; and the highest fitness, where the conserved structure in the RNA is present the highest faces, where the conserved structure in the RNA is present conserved structure in a RNA sequence. The present sequence is used in the exemplification of the present invention.
                                                                                                                                                                                                                                                          o;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       gulatory element-binding transcription factor, SRBBF, disorder; diabetes; cardiovascular disorder; atherosclerosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /mod_base= OTHER
/note= "Phosphorothioate linkages. All cytidines are 5-
                                                                                                                                                                                                                                                         Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               element-binding protein-1; SREBP-1; ss; mouse;
                                                                                                                                                                                                                                                         ·,
                                                                                                                                                                                                                             Length 20;
                                                                                                                                                                                                                           Score 13.8; DB 1; Length 2
Pred. No. 1e+03;
0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mouse SREBP-1 antisense oligonucleotide ISIS 219655.
                                                                                                                                                                                                Sequence 20 BP; 4 A; 6 C; 8 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /mod_base= OTHER
/note= "2'-methoxyethtyl residues'
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                'note= "2'-methoxyethyl residues"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Location/Qualifiers
             Example 1; Fig 11; 52pp; English.
                                                                                                                                                                                                                                                                               1239
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     methylcytidines"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Dobie KW;
                                                                                                                                                                                                                                                                                                         18
                                                                                                                                                                                                                                                                                                                                                                      ADG72117 standard; DNA; 20 BP
                                                                                                                                                                                                                         0.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    04-JUN-2002; 2002US-00161996
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                                                                                                                                                                                                                                                                              1223 TGGAGGAACAGCTACAC
                                                                                                                                                                                                                                                                                                     TGGAGGAGCAGCTCCAC
                                                                                                                                                                                                                                                                                                                                                                                                                          (first entry)
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                                                                                                                                                                                                                                       Local Similarity 88.2
1es 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      16. .20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           gene therapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  *tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /*tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /*tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Baker BF,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2004-022079/02
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                regulatory
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          regulatory
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    hyperlipidaemia.
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modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   modified base
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                                                                                                                                                                                                                           Query Match
                                                                                                                                                                                                                                                                                                                                                                                                 ADG72117;
                                                                                                                                                                                                                                       Best Local
Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sterol
                                                                                                                                                                                                                                                                                                                                             RESULT 1588
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         sterol
                                                                                                                                                                                                                                                                                                                                                           ADG72117
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The invention relates to a compound 8-80 nucleobases in length targeted to, and which specifically hybridises with a nucleic acid molecule encoding sterol regulatory element-binding protein-1 (SREBP-1, also known eas sterol regulatory element-binding protein-1 (SREBP-1, also known control regulatory element-binding protein-1 (SREBP-1, and control regulatory element-binding control control control control east an 8-nucleobase in length that specifically hybridises with at least an 8-nucleobase portion of an active site on a nucleic acid molecule encoding sterol regulatory element-binding protein-1, a composition comprising the compound and a carrier or diluent, or neells or tissues (by contacting the colls or tissues with the compound in cells or tissues (by contacting the cells or tissues with the compound contact with sterol regulatory element-binding protein-1 contacting the colls or tissues with the compound contacting an animal having a disease or condition companies of sterol regulatory element-binding protein-1 compound so that expression of sterol regulatory element-binding protein-1 compound so that expression of sterol regulatory element-binding protein-1 compound so that expression of sterol regulatory element-binding protein-1 compound so that expression of sterol regulatory element-binding protein-1 is almibited. The animal a therapeutic or prophylactic amount of the compound so that expression of sterol regulatory element-binding protein-1 is modified internucleoside linkage (preferably a phosphorothicate linkage), enter least one modified mucleobase (preferably 5'-O-methoxyethyl sugar modety) and modified mucleobase (preferably 5'-O-methoxyethyl for modified modified mucleobase (preferably 5'-O-methoxyethyl for an encodified modified mucleobase (preferably 5'-O-methoxyethyl for an encodified modified mucleobase (preferably 5'-O-methoxyethyl for an encodified modified mucleobase (preferably 5'-O-methoxyethyl sugar modified mucleobase (preferably 5'-O-methoxyethyl sugar modified mucleobase (preferab
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   -binding protein-1, such as a metabolic disorder e.g. diabetes, or a cardiovascular disorder, e.g. atherosclerosis or hyperlipidaemia. They are also useful in research and diagnostics for modulating the expression of sterol regulatory element-binding protein-1. The present sequence is an antisense oligonucleotide targeting mouse SREBP-1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  methylcytosine). The compound, composition and methods are useful for treating a disease or condition associated with sterol regulatory element
                   nucleic acid encoding sterol regulatory element-binding protein-1, useful for treating diabetes, atherosclerosis or hyperlipidemia.
particularly antisense oligonucleotides targeted to a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0.8%; Score 13.8; DB 1; Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 20 BP; 5 A; 8 C; 4 G; 3 T; 0 U; 0 Other;
                                                                                                           Example 16; SEQ ID NO 112; 112pp; English
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADG72234 standard; cDNA; 20 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  04-JUN-2002; 2002US-00161996.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mouse SREBP-1 target site #3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     88.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   20 regrescherence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Best Local Similarity
Matches 15; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     hyperlipidaemia
  New compounds,
nucleic acid er
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mus musculus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            11-MAR-2004
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADG72234;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RESULT 1589
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         sterol
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/note= "Phosphorothioate linkages. All cytidines are 5-methylcytidines"
                                                                                                                                                                                                                                                                                                                            sterol regulatory element-binding transcription factor; SREBF; metabolic disorder; atherosclerosis;
                                                                                                                                                                                                                                                                                                  Sterol regulatory element-binding protein-1; SREBP-1; ss; human;
                             0.8%; Score 13.8; DB 1; Length 20;
88.2%; Pred. No. 1e+03;
ive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                         Human SREBP-1 antisense oligonucleotide ISIS 220046.
                                                                                                                                                                                                                                                                                                                                                                                                                   Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                              /mod base= OTHER
                                                                                          1729 CACCTGCCCACTTGTCC 1745
                                                                                                                                                                                          BP.
                                                                                                                      CACCTGCACCCTTGTCC
                                                                                                                                                                                           ADG72049 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                Д
                                                                                                                                                                                                                                                (first entry)
                                                  Local Similarity 88.2
nes 15; Conservative
                                                                                                                                                                                                                                                                                                                      antisense gene therapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                    ..20
*tag=
                                                                                                                                                                                                                                                                                                                                                               hyperlipidaemia.
                                                                                                                                                                                                                                                                                                                                                                                                                       Key
modified_base
                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens.
                                                                                                                                                                                                                                                  11-MAR-2004
                                                                                                                                                                                                                       ADG72049;
                                   Query Match
                                                                                                                                                                 RESULT 1590
                                                                  Matches
                                                                                                                                                                               ADG72049
                                                     Best
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             ö
                                                                                                                                                                                                                                                                            regulatory element-binding transcription factor; SREBF;
lic disorder; diabetes; cardiovascular disorder; atherosclerosis;
              Gaps
                                                                                                                                                                                                                                                     Sterol regulatory element-binding protein-1; SREBP-1; ss; mouse; antisense gene therapy;
             ;
             Indels
Pred. No. 1e+03;
; Mismatches 2;
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(ISIS-) ISIS PHARM INC.

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Gaps

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The invention relates to a compound 8-80 mucleobases in length targeted to, and which specifically hybridises with a nucleic acid molecule encoding sterol regulatory element-binding protein-1 (SRBB-1, also known as sterol regulatory element-binding protein-1 (SRBB-1, also known as sterol regulatory element-binding pranscription factor, SRBBF), and inhibits the expression of SRBP-1, i.e. is an antisense oligonuclectide. Also included are a compound 8-80 mucleobase in length that specifically comparing are are a compound 8-80 mucleobase portion of an active site on a nucleic acid molecule encoding sterol regulatory element-binding protein-1 in composition comprising the compound and a carrier or diluent, inhibiting the expression of sterol regulatory element-binding protein-1 in compound so trissues (by contacting the cells or tissues with the compound computed and treating an animal having a disease or condition associated with sterol regulatory element-binding protein-1 by associated with sterol regulatory element-binding protein-1 is inhibited. The antisense oligonuclectide comprises at least one conditied sucar moiety (preferably 2'-0-methoxyethyl sugar moiety) or at least one modified sucar moiety (preferably 2'-0-methoxyethyl sugar moiety) or at least one modified nucleobase (preferably 5-methoxyethyl sugar moiety) or at least one modified nucleobase (preferably 5-methoxyethyl sugar creating a disease or condition associated with sterol regulatory element-binding protein-1, such as a metabolic disorder e.g. diabetes, or are also useful in research and diagnostics for modulating the expression of sterol regulatory element-binding protein-1. The present sequence is a mean are also useful for the antisense oligonucleotides.
                                                                                                 New compounds, particularly antisense oligonuclectides targeted to a nucleic acid encoding sterol regulatory element-binding protein-1, useful for treating diabetes, atherosclerosis or hyperlipidemia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 20 BP; 3 A; 9 C; 3 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                       Example 16; SEQ ID NO 229; 112pp; English.
  Dobie KW;
Baker BF,
                                                      WPI; 2004-022079/02.
     Freier SM,
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13:39:09 2004

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Tue Nov

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New compounds, particularly antisense oligonucleotides targeted to a nucleic acid encoding sterol regulatory element-binding protein-1, useful for treating diabetes, atherosclerosis or hyperlipidemia.
                 /note= "2'-methoxyethyl residues"
                                                                                                                                                              Example 15; SEQ ID NO 44; 112pp; English.
            /mod_base= OTHER
                                                                            04-JUN-2002; 2002US-00161996
                                                                                       04-JUN-2002; 2002US-00161996
1.
/*tag=
'*
                                                                                                    (ISIS-) ISIS PHARM INC
                                                                                                               Freier SM, Baker BF,
                                                                                                                           WPI; 2004-022079/02.
                                                    US2003224515-A1
modified base
                       modified base
                                                                04-DEC-2003
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The invention relates to a compound 8-80 nucleobases in length targeted co, and which specifically hybridises with a nucleic acid molecule conciding sterol regulatory element-binding protein-1 (SRBP-1, also known as sterol regulatory element-binding protein-1 (SRBP-1, also known in the capture are a compound 8-80 nucleobases in length that specifically hybridises with at least an 8-nucleobase sin length that specifically hybridises with a least an 8-nucleobase portion of an active site on a nucleic acid molecule encoding sterol regulatory element-binding protein-1 inhibiting the expression of sterol regulatory element-binding protein-1 inhibiting the expression of sterol regulatory element-binding protein-1 in cells or tissues (by contacting the cempound and acarrier or diluent, inhibited) and treating an animal having a disease or condition associated with sterol regulatory element-binding protein-1 is inhibited and treating a natural having a disease or condition compound so that expression of sterol regulatory element-binding protein-1 is inhibited. The antisense oligonucleotide comprises at least one modified sugar moiety (preferably 2, -0-methoxyethyl sugar moiety) or at least one modified sugar moiety (preferably 2, -0-methoxyethyl sugar moiety) or at least one modified sugar moiety (preferably 2, -0-methoxyethyl sugar moiety) or at least one modified sugar moiety (preferably 2, -0-methoxyethyl sugar moiety) or at least one modified sugar moiety (preferably 2, -0-methoxyethyl sugar cardiovascular disorder, e.g. atherosclarosis or hyperlipidaemia. They are also useful in research and diagnostics for modulating the expression of sterol regulatory element-binding protein-1. The present sequence is an antisense oligonucleotide targeting human SREBP-1.

Sequence 20 BP; 5 A; 4 C; 7 G; 4 T; 0 U; 0 Other;

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·,
     DB 1; Length 20;
                                2; Indels
  0.8%; Score 13.8; DB 1
88.2%; Pred. No. 1e+03;
ative 0; Mismatches
                                                                                    18
                                                                                 GAGGCTTCCTGCAGAA
                                                      505 GAGGGCTACCTGGAGAA
Query Match 0.8
Best Local Similarity 88.2
Matches 15; Conservative
                                                      ð
                                                                              g
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ADG72186 standard; DNA; 20 RESULT 1591

ADG72186; ADG72186/c ID ADG72 XX ADG72

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Human SREBP-1 target site #23
11-MAR-2004 (first entry)
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sterol regulatory element-binding transcription factor; SREBF; metabolic disorder; diabetes; cardiovascular disorder; atherosclerosis; element-binding protein-1; SREBP-1; ds; human; antisense gene therapy; regulatory hyperlipidaemia. Sterol

Homo sapiens.

US2003224515-A1.

04-DEC-2003

04-JUN-2002; 2002US-00161996

04-JUN-2002; 2002US-00161996

(ISIS-) ISIS PHARM INC

Dobie KW; Freier SM, Baker BF,

WPI; 2004-022079/02.

New compounds, particularly antisense oligonucleotides targeted to a nucleic acid encoding sterol regulatory element-binding protein-1, useful for treating diabetes, atherosclerosis or hyperlipidemia.

Example 16; SEQ ID NO 181; 112pp; English

The invention relates to a compound 8-80 nucleobases in length targeted to, and which specifically hybridises with a nucleic acid molecule concoling sterool regulatory element-binding protein-1 (SREBP-1, also known as sterool regulatory element-binding transcription factor, SREBP), and inhibits the expression of SREBP-1, i.e. is an antisense oligonucleotide. Also included are a compound 8-80 nucleobases in length that specifically conversely with at least an 8-nucleobases in length that specifically succession of sterol regulatory element-binding protein-1, a composition comprising the compound and a carrier or diluent, in cells or tissues (by contacting the cells or tissues with the compound of sterol regulatory element-binding protein-1 in cells or tissues (by contacting the cells or tissues with the compound of sterol regulatory element-binding protein-1 is associated with sterol regulatory element-binding protein-1 by associated with sterol regulatory element-binding protein-1 is associated with sterol regulatory (preferably a phosphorothicate linkage), at least one modified supar moiety (preferably 2'-O-methoxyethyl sugar moiety) or at least one modified supar moiety (preferably 2'-O-methoxyethyl sugar moiety) or at least one modified nucleobase (preferably 5-moiety) creating a disease or condition associated with sterol regulatory element compound, composition and methods are useful for treating a disease or condition associated with sterol regulatory element condition associated with sterol regulatory element of attached are also useful in research and diagnostics for modulating the expression of sterol regulatory element ending a prisearch and insease oligonucleotice is a human SRBP-1 target region for the arrisense oligonucleotice is a longer region for the arrisense oligoner is a present sequence is a human SREBP-1 target region for the antisense oligonucleotides

Sequence 20 BP; 4 A; 7 C; 4 G; 5 T; 0 U; 0 Other;

ó:

Gaps

Gaps . 0 DB 1; Length 20; 2; Indels 0.8%; Score 13.8; DB 1 18.2%; Pred. No. 1e+03; ve 0; Mismatches 88.2%; 15; Conservative Local Similarity Query Match Matches

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505 GAGGCTACCTGGAGAA 521 19 GAGGGCTTCCTGCAGAA 3 ð 셤

RESULT 1592

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treating a disease or condition associated with sterol regulatory element
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New compounds, particularly antisense oligonucleotides targeted to a nucleic acid encoding sterol regulatory element-binding protein-1, useful for treating diabetes, atherosclerosis or hyperlipidemia.
                                                                                                                                                                                                                                                                                                        antisense gene therapy;
sterol regulatory element-binding transcription factor; SREBF;
metabolic disorder; diabetes; cardiovascular disorder; atherosclerosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /note= "Phosphorothioate linkages. All cytidines are 5-
                                                                                                                                                                                                                                                                   regulatory element-binding protein-1; SREBP-1; ss; mouse;
                                                                                                                                                                                                                Mouse SREBP-1 antisense oligonucleotide ISIS 219640.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             "2'-methoxyethtyl residues"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example 16; SEQ ID NO 105; 112pp; English.
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/*tag= c
/mod base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               methylcytidines"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         'mod_base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /mod_base= OTHER
/note= "2'-metho
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Dobie KW;
                                                           BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   04-JUN-2002; 2002US-00161996
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                                                           ADG72110 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Д
                                                                                                                                                                      (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Baker BF,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2004-022079/02.
                                                                                                                                                                                                                                                                                                                                                                                       hyperlipidaemia.
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modified base
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                                                                                                                                                                                                                                                                                                                                                                                                                                               Mus musculus.
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                                                                                                                                                                         11-MAR-2004
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                                                                                                                   ADG72110;
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                                                                                                                                                                                                                                                                                       Sterol
                                    ADG72110/c
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inhibited and treating an animal having a disease or condition associated and treating an animal having a disease or condition associated with sterol regulatory element-binding protein-1 by administering to the animal a therapeutic or prophylactic amount of the compound so that expression of sterol regulatory element-binding protein-1 is inhibited. The antisense oligonucleotide comprises at least one modified internucleoside linkage (preferably a phosphorotothioare linkage), an least one modified sugar moiety (preferably 2.0-methoxyethyl sugar moiety) or at least one modified nucleobase (preferably 5-methoxyethyl sugar methylcytosine). The compound, composition and methods are useful for

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The invention relates to a compound 8-80 nucleobases in length targeted cto, and which specifically hybridises with a nucleic acid molecule cto, and which specifically hybridises with a nucleic acid molecule encoding sterol regulatory element-binding transcription factor, SREBF), and as sterol regulatory element-binding transcription factor, SREBF), and cto included are a compound 8-80 nucleobases in length that specifically hybridises with at least an 8-nucleobases in length that specifically cucleic acid molecule encoding sterol regulatory element-binding protein-ctor inhibiting the expression of sterol regulatory element-binding protein-ctor includes of sterol regulatory element-binding protein-ctor includes of sterol regulatory element-binding protein-ctor included and treating an animal having a disease or condition associated with sterol regulatory element-binding protein-lise associated with sterol regulatory element-binding protein-lise compound so that expression of sterol regulatory element-binding protein-lise administering to the animal a therapeutic or prophylactic amount of the compound so that expression of sterol regulatory element-binding protein-compound so that expression element-binding protein-compound 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New compounds, particularly antisense oligonucleotides targeted to a nucleic acid encoding sterol regulatory element-binding protein-1, useful for treating diabetes, atherosclerosis or hyperlipidemia.
                                                                                                                                                                                                                                                                                                           ö
-binding protein-1, such as a metabolic disorder e.g. diabetes, or a cardiovascular disorder, e.g. atherosclerosis or hyperlipidaemia. They are also useful in research and diagnostics for modulating the expression of sterol regulatory element-binding protein-1. The present sequence is an antisense oligonucleotide targeting mouse SREBP-1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  sterol regulatory element-binding transcription factor; SREBF; metabolic disorder; diabetes; cardiovascular disorder; atherosclerosis;
                                                                                                                                                                                                                                                                                                              Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sterol regulatory element-binding protein-1; SREBP-1; ss; mouse;
                                                                                                                                                                                                                                                                                                              . 0
                                                                                                                                                                                                                                            0.8%; Score 13.8; DB 1; Length 20;
88.2%; Pred. No. 1e+03;
tive 0; Mismatches 2; Indels
                                                                                                                                                                                           Sequence 20 BP; 5 A; 3 C; 9 G; 3 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                            1729 CACCTGCCCACTTGTCC 1745
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADG72241 standard; cDNA; 20 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mouse SREBP-1 target site #10.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 04-JUN-2002; 2002US-00161996.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
                                                                                                                                                                                                                                                                                                                       15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         antisense gene therapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (ISIS-) ISIS PHARM INC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2004-022079/02
                                                                                                                                                                                                                                                                                           Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          hyperlipidaemia.
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                                                                                                                                                                                                                                                            Query Match
Best Local S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        RESULT 1593
                                                                                                                                                                                                                                                                                                                             Matches
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            at least one modified sugar mojety (preferably 2-0-methoxyethyl sugar moiety) or at least one modified nucleobase (preferably 5-0-methoxyethyl sugar methylcytosine). The compound, composition and methods are useful for treating a disease or condition associated with sterol regulatory element binding protein-1, such as a metabolic disorder e.g. diabetes, or a cardiovascular disorder, e.g. atherosclerosis or hyperlipidemia. They are also useful in research and diagnostics for modulating the expression of sterol regulatory element-binding protein-1. The present sequence is a
                                                                                                                                                                                                                        ·,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The invention comprises an antisense oligonucleotides that are targeted to nucleic acids encoding a mammalian glucocorticoid receptor. The antisense oligonucleotides of the invention are useful for preventing or delaying infection, inflammation or tumour formation. The antisense oligonucleotides are also useful for treating diabetes, obesity, cardiovascular disorders, hyperlipidaemia or Cushing's syndrome. The present DNA sequence represents an antisense oligonucleotide that targets the human glucocorticoid receptor gene. NOTE: The present sequence contains 2'-methoxyethyl (2'-MOE) wings and a phosphorothioate backbone.
 modified internucleoside linkage (preferably a phosphorothioate linkage),
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           mammalian glucocorticoid receptor, useful for treating diabetes, obesity, cardiovascular disorder, hyperlipidemia or Cushing's syndrome.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     antisense oligonucleotide; glucocorticoid receptor; infection; inflammation; tumour formation; diabetees; obesity; cardiovascular disorder; hyperlipidaemia; Cushing's syndrome; human; ss; phosphorothioate backbone; 2-methoxyethyl; 2'-MOE.
                                                                                                                                                                                                                                                                                                                                                                                                                                         Human glucocorticoid receptor-specific antisense oligonucleotide #4116.
                                                                                                                                                                                                                        Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New antisense compound targeted to a nucleic acid molecule encoding
                                                                                                                                mouse SREBP-1 target region for the antisense oligonucleotides
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0
                                                                                                                                                                                           20;
                                                                                                                                                                                                                       2; Indels
                                                                                                                                                                                        Score 13.8; DB 1; Length
Pred. No. 1e+03;
0; Mismatches 2; Indels
                                                                                                                                                               Sequence 20 BP; 3 A; 4 C; 8 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 20 BP; 5 A; 5 C; 8 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 4; SEQ ID NO 4116; 985pp; English
                                                                                                                                                                                                                       0
                                                                                                                                                                                                                                                  240 TGGCGCCAGTGACCCTG 256
                                                                                                                                                                                                                                                                                                                                           282/c
ADH67282 standard; DNA; 20 BP.
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                                                                                                                                                                                          0.8%;
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                                                                                                                                                                                                                                                                                                                                                                                                            25-MAR-2004 (first entry)
                                                                                                                                                                                        Query Match 0.8
Best Local Similarity 88.2
Matches 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Nalseth AE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                 ADH67282
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ADH67282/c
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DB 1; Length 20;

Score 13.8; DB Pred. No. 1e+03;

0.8**%**; 88.2**%**;

Best Local Similarity

Query Match

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0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New antisense compound targeted to a nucleic acid molecule encoding mammalian glucocorticoid receptor, useful for treating diabetes, obesity, cardiovascular disorder, hyperlipidemia or Cushing's syndrome.
                                                                                                                                                                                                                                                                                                                                                                                                                             antisense oligonucleotide; glucocorticoid receptor; infection; inflammation; tumour formation; diabetes; obesity; cardiovascular disorder; hyperlipidaemia; Cushing's syndrome; human; ss; phosphorothioate backbone; 2'-methoxyethyl; 2'-MOE.
                                                                                                                                                                                                                                                                                                                                                                                  Human glucocorticoid receptor-specific antisense oligonucleotide #3762.
     Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
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  Indels
  7
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Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 4; SEQ ID NO 3762; 985pp; English.
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0
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                                               49 CCAGCAGTGTGACTGCT
                                                                                               18 ccascasrerecriecr
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                                                                                                                                                                                                                                                                                                                             (first entry)
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15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Similarity 88.2
15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Crosby SD, Nalseth AE;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (PHAA ) PHARMACIA CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2004-035034/03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WO2003099215-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Homo sapiens.
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Best Local 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           49
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Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Matches
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ID ADH54
XX AC ADH54
XX DT 25-M5
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The invention relates to an antisense oligonucleotide targeted to a nucleic acid encoding Livin and that specifically hybridises with the nucleic acid encoding Livin and inhibits expression of Livin. The antisense oligonucleotide is useful for preparing a composition for treating hyperproliferative disorder or aberrant apoptosis. The present sequence represents a human livin antisense oligonucleotide.
                                                                                                                                                                                                       The invention relates to an antisense oligonucleotide targeted to a nucleic acid encoding Livin and that specifically hybridises with the nucleic acid encoding Livin and inhibits expression of Livin. The antisense oligonucleotide is useful for preparing a composition for treating hyperproliferative disorder or aberrant apoptosis. The present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New antisense oligonucleotide, having a sequence targeted to a nucleic acid encoding Livin, useful for preparing a composition for treating hyperproliferative disorder or aberrant apoptosis.
                                                                                       New antisense oligonucleotide, having a sequence targeted to a nucleic acid encoding Livin, useful for preparing a composition for treating hyperproliferative disorder or aberrant apoptosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           hyperproliferative disorder; aberrant apoptosis; human; ss; Livin;
                                                                                                                                                                                                                                                                                                                                                                                           DB 1; Length 20
                                                                                                                                                                                                                                                                                                                                                                                                                                  2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human Livin antisense oligonucleotide ISIS 205823.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 20 BP; 4 A; 9 C; 5 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                     Sequence 20 BP; 2 A; 5 C; 9 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                              sequence represents a human livin target region.
                                                                                                                                                                                                                                                                                                                                                                                        Score 13.8; DB 1
Pred. No. 1e+03;
0; Mismatches
                                                                                                                                                                          Example 15; SEQ ID NO 116; 60pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 15; SEQ ID NO 41; 60pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1645
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   02-JUL-2002; 2002US-00188646.
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                                                                                                                                                                                                                                                                                                                                                                                               0.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1629 CCCCAGCAGCAGCGGC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CCTCAGCAGTCAGCGGC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first entry)
                                                                                                                                                                                                                                                                                                                                                                                           Query Match 0.8
Best Local Similarity 88.2
Matches 15, Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Dobie KW;
                Dobie KW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2004-098436/10.
                                                 WPI; 2004-098436/10.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  22-APR-2004
                Bennett CF,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADH89572
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RESULT 1598
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Ношо
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADH89572
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The invention relates to a compound targeted to and which specifically hybridises with a nucleic acid molecule encoding VBGF-C, and inhibits the expression of VBGF-C. The compound, composition and methods are useful for treating a disease or condition associated with VBGF-C, such as a cardiovascular disorder e.g. atherosclerosis or diabetic retinopathy or an autoimmune or inflammatory disorder. They are also useful in research and diagnostics for modulating the expression of VBGF-C. The present sequence represents a human VBGF-C antisense oligomucleotide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                stacid encoding vascular endothelial growth factor-C (VBGF-C), for treating atherosclerosis, diabetic retinopathy, or
                                                                                                                                                                                                                                                                                                                                                                                                                                                             particularly antisense oligonucleotides targeted to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              hyperproliferative disorder; aberrant apoptosis; human; ss; Livin.
                                   human; ss; VEGF-C; cardiovascular disorder; atherosclerosis; diabetic retinopathy; autoimmune disorder; inflammatory disorder; vascular endothelial growth factor; antisense.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0.8%; Score 13.8; DB 1; Length 20; 88.2%; Pred. No. 1e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Indels
Human VEGF-C antisense oligonucleotide ISIS 158128
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 20 BP; 5 A; 6 C; 4 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example 15; SEQ ID NO 41; 83pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human Livin target region ISIS 123483
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1564 ATGCCTGACTCAGGCAG 1580
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    BP
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                                                                                                                                                                                                                                                                17-JUN-2002; 2002US-00173718
                                                                                                                                                                                                                                                                                                     17-JUN-2002; 2002US-00173718
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          inflammatory disorders.
                                                                                                                                                                                                                                                                                                                                             (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2004-061284/06.
                                                                                                                                                                                                                                                                                                                                                                                  Dobie KW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity
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                                                                                                                                                                                   US2003232437-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   compounds,
                                                                                                                                             Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 22-APR-2004
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                                                                                                                                                                                                                          18-DEC-2003
                                                                                                                          Synthetic
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RESULT 1597

Matches

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Homo

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Gaps

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Zhang

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Gaps

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10017621-381.rng

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The invention relates to novel compounds of 8-80 nucleobases in length targeted to, and which specifically hybridises with, a nucleic acid molecule encoding 3-hydroxy-3-methylglutary1-Coenzyme A (EMG-CoA) reductase, and inhibits the expression of HMG-COA reductase. The novel compounds have cardiant, antiarteriosclerotic, and antilipaemic activities. The compound can be used to treat disorders by antisense gene therapy. The compounds, compositions and methods are useful for treating a disease or condition associated with HMG-CoA reductase, such as a cardiovascular disorder e.g. atherosclerosis, or a disease or condition involving cholesterol metabolism. They are also useful in research and diagnostics for modulating the expression of HMG-CoA reductase. This polynucleotide sequence represents an antisense oligonucleotide of the
                                                                                                                                                                                                                                                                                                           HMG-COA reductase; 3-hydroxy-3-methylglutaryl-Coenzyme A;
HMG-COA reductase; cardiant; antiarteriosclerotic; antilipaemic;
antisense gene therapy; cardiovascular disorder; cholesterol metabolism;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New compounds, particularly antisense oligonucleotides targeted to
nucleic acid encoding HMG-CoA reductase, useful for treating
atherosclerosis, or a disease involving cholesterol metabolism or
                                                                                                                                                                                                                                                                             Human HMG-CoA reductase antisense oligonucleotide, SEQ ID No 100
   ; DB 1; Length 20;
Le+03;
                                     Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 20 BP; 7 A; 10 C; 1 G; 2 T; 0 U; 0 Other;
                                   2;
   Score 13.8; D
Pred. No. 1e+0
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example 15; SEQ ID NO 100; 110pp; English.
                                                                1629 CCCCAGCAGCAGCGGC 1645
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Dobie KW;
                                                                                                 13
                                                                                                                                                                                 BP
0.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             02-JUL-2002; 2002US-00190366.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           02-JUL-2002; 2002US-00190366.
                                                                                             3 cercadeagreagege
                                                                                                                                                                            ADI79577 standard; DNA; 20
                                                                                                                                                                                                                                            (first entry)
                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Freier SM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 2004-081743/08.
                 Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                            US2004006031-A1.
                                                                                                                                                                                                                                                                                                                                                                                               Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              angiogenesis.
                                                                                                                                                                                                                                              22-APR-2004
                                15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                            08-JAN-2004
                                                                                                                                                                                                                                                                                                                                                             human; ss.
                                                                                                                                                                                                              ADI79577;
   Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Dean NM,
                                                                                                                                                RESULT 1599
                                Matches
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The invention relates to novel compounds of 8-80 nucleobases in length targeted to, and which specifically hybridises with, a nucleic acid molecule encoding 3-hydroxy-3-methylglutary1-Conzyme A (HMG-COA) reductase, and inhibits the expression of HMG-COA reductase. The novel compound can be used to treat disorders by antichsense therapy. The compound can be used to treat disorders by antichsense therapy. The compounds, compositions and methods are useful for treating a disease or condition associated with HMG-COA reductase, such as a
                                                                                                                         HWG-CoA reductase, 3-hydroxy-3-methylglutaryl-Coenzyme A;
HMG-CoA reductase; cardiant; antiarteriosclerotic; antilipaemic;
antisense gene therapy; cardiovascular disorder; cholesterol metabolism;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        cardiovascular disorder e.g. atherosclerosis, or a disease or condition involving cholesterol metabolism. They are also useful in research and diagnostics for modulating the expression of HMG-CoA reductase. This
                                                                                                                                                                                                                                                                                                                                                                                                                                                              ø
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                      New compounds, particularly antisense oligonucleotides targeted to
nucleic acid encoding HMG-CoA reductase, useful for treating
atherosclerosis, or a disease involving cholesterol metabolism or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     polynucleotide sequence represents an antisense oligonucleotide of
                                                                                            Human HMG-CoA reductase antisense oligonucleotide, SEQ ID No 297.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 13.8; DB 1; Length 20; Pred. No. 1e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human LIM domain kinase 1 antisense oligonucleotide #104.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 20 BP; 2 A; 1 C; 10 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example 16; SEQ ID NO 297; 110pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           BP.
                                                                                                                                                                                                                                                                                                                              02-JUL-2002; 2002US-00190366
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                                                                                                                                                                                                                                                                                                                                                                                             Dobie
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0.88;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                88.28;
 ADI79774 standard; DNA; 20
                                                               (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               465 CAACAAGCGCCTATCAC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CAACAAGCTCCCATCAC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADI38820 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                             (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                          Dean NM, Freier SM,
                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2004-081743/08.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Local Similarity
                                                                                                                                                                                                                                    US2004006031-A1.
                                                                                                                                                                                                      Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      angiogenesis.
                                                               22-APR-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     22-APR-2004
                                                                                                                                                                                                                                                                 08-JAN-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   15;
                                                                                                                                                                         human; ss.
                                ADI 79774;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADI38820;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RESULT 1601
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADI38820
ID ADI3
XX ADI3
XX DT 22-7
XX DE HUM?
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Gaps

0;

2; Indels

Pred. No. 1e+03; ; Mismatches

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Conservative

88.2%;

Query Match Best Local Similarity Matches 15; Conserv

CAACAAGCGCCTATCAC 481

465

à

CAACAAGCTCCCATCAC 19

0.8%; Score 13.8;

DB 1; Length 20;

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WPI; 2004-121553/12.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Similarity
                                                                                                                                                                                                                        US2004014047-A1
                                                               Key
modified_base
                                                                                                                           modified_base
                                                                                                                                                                    modified base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           22-APR-2004
                                           Homo sapiens
                                                                                                                                                                                                                                                                                                                                Cowsert LM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       15;
                                                                                                                                                                                                                                             22-JAN-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADI38608;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              RESULT 1603
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Matches
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1D AD13

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AC AD13

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DT 22-A
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                                                                                                                                                                                                                                                                                                                                                                                                                           The invention describes a compound 8-80 nucleobases in length targeted to a nucleic acid molecule encoding LIM domain kinase 1. The compound specifically hybridises with the nucleic acid molecule encoding LIM domain kinase 1 and inhibits the expression of LIM domain kinase 1. It specifically hybridises with at least an 8-nucleobase portion of a preferred target region on the nucleic acid molecule encoding LIM domain kinase 1. The antisense oligonucleotide is useful for modulating the expression of LIM domain kinase 1 in cells or tissues to treat diseases associated with their expression, such as a developmental disorder or a neurological disorder. In addition, the compound is used for diagnosties.
                                                                                                                                                                                                                                                                                                                                                               New antisense oligonucleotides for modulating LIM domain kinase 1 expression, useful for diagnosing, preventing or treating conditions associated with the kinase, e.g. neurological or developmental disorders.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     prophylăxis, or as research reagents or kits. This sequence represents a human LIM domain kinase 1 antisense oligonucleotide.
                                                                                  /mod_base= OTHER
/note= "OTHER= Phosphorothioate backbone. All cytidines
are 5-methylcytidines"
1. .5
                                                                                                                                                                          *tag= a
|mod_base= OTHER
'note= "OTHER= 2'-0-Methoxyethyl (2'-MOE) nucleotides"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0;
  neuroprotective; LIM domain kinase 1; developmental disorder; neurological disorder; diagnostic; prophylaxis; human; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0.8%; Score 13.8; DB 1; Length 20; 88.2%; Pred. No. 1e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human LIM domain kinase 1 antisense oligonucleotide #33.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 20 BP; 4 A; 9 C; 5 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                             Example 15; SEQ ID NO 119; 81pp; English
                                                        Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    760 TCCCTGCTCAAGGACCT 776
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADI38749 standard; DNA; 20 BP
                                                                                                                                                                                                                                                            18-JUL-2002; 2002US-00199199
                                                                                                                                                                                                                                                                                 18-JUL-2002; 2002US-00199199
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      4 TCCCAGCGCAAGGACCT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     22-APR-2004 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  15; Conservative
                                                                                                                                                               15. .20
                                                                                                                                                                                                                                                                                                      (ISIS-) ISIS PHARM INC.
                                                                            *tag=
                                                                    . 20
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                                                                                                                                                                                                                                                                                                                                               WPI; 2004-121553/12.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity
Matches 15; Conserv
                                                                                                                                                                                                                    US2004014047-A1
                                                          Key
modified_base
                                                                                                                        modified base
                                                                                                                                                                modified base
                                      sapiens
                                                                                                                                                                                                                                         22-JAN-2004
                                                                                                                                                                                                                                                                                                                           Cowsert LM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADI38749;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RESULT 1602
                                      Homo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADI38749
     BXAXAXB
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New antisense oligonucleotides for modulating LIM domain kinase 1 expression, useful for diagnosing, preventing or treating conditions associated with the kinase, e.g. neurological or developmental disorders.
                                                                                                                                                                                                                                          /mod_base= OTHER /note= "OTHER= Phosphorothioate backbone. All cytidines are 5-methylcytidines"
                                                                                                                                                                                                                                                                                                                                                                                                      /*tag= a
/mod_base= OTHER
/note= "OTHER= 2'-0-Methoxyethyl (2'-MOE) nucleotides"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /note= "OTHER= 2'-O-Methoxyethyl (2'-MOE) nucleotides"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ·
0
neuroprotective; LIM domain kinase 1; developmental disorder; neurological disorder; diagnostic; prophylaxis; human; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 20 BP; 2 A; 5 C; 9 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example 15; SEQ ID NO 48; 81pp; English.
                                                                                                                                                                  socation/Qualifiers
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/mod base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     776
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADI38608 standard; DNA; 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              15. .20
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                                                                                                                                                                                                               1. .20
/*tag=
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Phosphorothioate backbone. All cytidines

are 5-methylcytidines"

/*tag= a

modified base

/mod_base= OTHER 'note= "OTHER=

Д

...20 *tag=

modified_base

sapiens

Homo

/mod_base= OTHER /note= "OTHER= Phosphorothioate backbone. All cytidines are 5-methylcytidines"

Location/Qualifiers

Д

/*tag=

/mod_base= OTHER /note= "OTHER= 2'-0-Methoxyethyl (2'-MOE) nucleotides"

/note= "OTHER= 2'-O-Methoxyethyl (2'-MOE) nucleotides

/mod base= OTHER

O

15. .20 /*tag=

*tag=

Location/Qualifiers

cytostatic; antidiabetic; antiinfertility; gene therapy; cyclin-dependent kinase 4; diabetes; infertility; hyperproliferative disorder; cancer; antisense technology; human;

Cyclin dependent kinase 4 antisense oligonucleotide #50

(first entry)

22-APR-2004

ADI26884;

ВP

ADI26884 standard; DNA; 20

/mod_base= OTHER /note= "OTHER= 2'-O-Methoxyethyl (2'-MOE) nucleotides" 15. .20

/mod_base= OTHER /note= "OTHER= 2'-0-Methoxyethyl (2'-MOE) nucleotides"

U

'*tag=

modified base

Dobie KW;

(ISIS-) ISIS PHARM INC Dean NM, Freier SM, WPI; 2004-081710/08.

02-JUL-2002; 2002US-00188779 02-JUL-2002; 2002US-00188779.

US2004005567-A1

08-JAN-2004

13:39:09 2004

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Tue Nov

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New antisense oligonucleotides for modulating dual specific phosphatase (expression, useful for diagnosing, preventing or treating conditions associated with the phosphatase, e.g. hyperproliferative or inflammatory
                cytostatic; antiinflammatory; antisense therapy; dual specific phosphatase 6; hyperproliferative disorder; apoptosis; inflammatory disorder; developmental disorder; diagnostic; prophylaxis; human; antisense technology; ss.
Dual specific phosphatase 6 antisense oligonucleotide #17
                                                                                                                                                                                                                                                                                                                                                                                          Example 15; SEQ ID NO 29; 54pp; English.
                                                                                                                                                                                                                                                      18-JUL-2002; 2002US-00199221
                                                                                                                                                                                                                                                                        18-JUL-2002; 2002US-00199221
                                                                                                                                                                                                                                                                                                            Cowsert LM,
                                                                                                                                                                                                                                                                                          (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                                                                                             WPI; 2004-121554/12.
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                                                                                        modified base
                                                                                                                                     modified base
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                                                               Homo sapiens
                                                                                                                                                                                                                                     22-JAN-2004
                                                                                                                                                                                                                                                                                                            Monia BP,
                                                                                                                                                                                                                                                                                                                                                                         disorders
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
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Dobie KW;

New antisense oligonucleotide, having a sequence targeted to a nucleic acid encoding cyclin-dependent kinase 4, useful for preparing a composition for treating diabetes, infertility or hyperproliferative disorder, e.g., cancer 19 Query Match Matches 셤 à The invention describes a compound 8-80 nucleobases in length targeted to a nucleic acid molecule encoding dual specific phosphatase 6. The compound specifically hybridises with the nucleic acid molecule encoding dual specific phosphatase 6 and inhibits the expression of dual specific phosphatase 6. It specifically hybridises with at least an 8-nucleobase portion of a preferred target region on the nucleic acid molecule concoming dual specific phosphatase 6. The antisense oligonucleotide is useful for inhibiting the expression of dual specific phosphatase 6 in cells or tissues to treat diseases associated with their expression, such a propriorilerative disorder, a condition arising from aberrant apoptosis, an inflammatory disorder or a developmental disorder. In addition, the compound is used for diagnostics, prophylaxis, or as research reagents or kits. This sequence represents a human dual specific phosphatase 6 antisense oligonucleotide. 0 Gaps o; 0.8%; Score 13.8; DB 1; Length 20; 18.2%; Pred. No. 1e+03; ve 0; Mismatches 2; Indels Indels Sequence 20 BP; 7 A; 4 C; 6 G; 3 T; 0 U; 0 Other; 2; 461 1 AAGATCTCCACTGGGAA 17 88.2%; 445 AAGATCTCCACTGAGGA Conservative Local Similarity nes 15; Conserv

705 0.8%; m 689 ACCTIGIGGCACTCAAG ACTITGEGCCCTCAAG Conservative Local Similarity 15;

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The invention describes a new antisense oligonucleotide, having a sequence comprising 8-80 bp targeted to a nucleic acid encoding cyclindependent kinase 4, specifically hybridises with the nucleic acid encoding cyclin-dependent kinase 4 and inhibits expression of cyclindependent kinase 4. The antisense oligonucleotide is useful for preparing a composition for treating diabetes, infertility or hyperproliferative disorder, e.g., cancer. This sequence represents a human cyclin dependent kinase 4 antisense oligonucleotide.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 13.8; DB 1; Length 20;
Pred. No. 1e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Indels
                                                                                                                                                                                                                                                                                                                                                                     Sequence 20 BP; 5 A; 5 C; 6 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0; Mismatches
Example 15; SEQ ID NO 69; 90pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADI79383 standard; DNA; 20
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BP.

RESULT 1604 ADI26884/c

Matches

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Sequence 20 BP; 2 A; 5 C; 6 G; 7 T; 0 U; 0 Other;

g

tumour progression; cytostatic; immunosuppressive; neuroprotective; cardiant; antimicrobial; nephrotropic; respiratory; vaccine; gene therapy; autoimmune disease; infectious disease; growth disturbance disease; neurological disease; cardiovascular disease; respiratory disease; kidney disease; neoplastic disease; mouse; Sema3C; semaphorin; reverse transcriptase; PCR; primer; ss. disease; disease pathogenesis; malignant; cancer; metastasis; Mouse Sema3C reverse transcriptase PCR primer SEQ ID NO:21. Albrechtsen M; Olsen O, 10-JUL-2003; 2003WO-DK000486 Lukanidin E, 11-JUL-2002; 2002DK-00001092 (first entry) WPI; 2004-122767/12. (SEMA-) SEMA APS WO2004006898-A2 Christensen C, 22-APR-2004 22-JAN-2004 Mus sp. Synthetic. ADI79383;

Use of an agent for preparing a medicament for preventing progression of an invasive disease, for treating malignant forms of cancer or for preventing metastasis of cancer in vivo and tumor progression in vitro and/or in vivo.

Example 1; SEQ ID NO 21; 134pp; English.

The present invention describes an agent which can be used for preparing a medicament for preventing the progression of an invasive disease in an individual, where invasion of cells, other organisms or invasion of cells intension of cancer in vivo and tumour of cancer or for preventing metastasis of cancer in vivo and tumour corporassion in vitro and/or in vivo. The agent comprises (a) an agent cambon in intro and/or in vivo. The agent comprises (a) an agent cambon in family of proteins; (b) an agent capable of inhibiting expression of a polypeptide belonging to the semaphorin family of proteins, where the agent is belonging to the semaphorin family of proteins, where the agent is selected from antibodies or fragments of antibodies directed to the polypeptide, or fragments or variants of fragments of the polypeptide, or fragments or variants of fragments of the polypeptide, or fragments or variants of fragments of the polypeptide, or fragment or a receptor and thereby inhibiting binding of a proteolytic and/or (c) an agent capable of inhibiting binding of a proteolytic convertase; (a) an isolated polyclonal activation of the convertase; (a) an isolated polyclonal or monoclonal antibody compound; or a proprotein convertase and thereby inhibiting the activity of the convertase; (3) an isolated polyclonal or monoclonal antibody compound; (5) a method for diagnosing or prognosing manignant cancer. The agent has expression of an invasive disease, where invasion of convertance, growth disturbance, neuroprotective, cardiant, antimicrobial, in gene therapy. The agent is useful for preparing a medicament for preventing progression of inself plays a role in disease.

Crespiratory, kidney or neoplastic diseases, where invasion of the present sequence represents a reverse transcriptase PCR primer crownerses; of cancer in vivo and tumour progression in vitro and/or in vention.

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                                                                                                                                                                                                                                                                                                                                                      Phosphorothicate backbone. All cytidines
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New antisense oligonucleotide, having a sequence targeted to a nucleic acid encoding PCTAIRE protein kinase 2, useful for preparing a composition for treating neurological disorders.
                                                                                                                                                                                                                                                                                                                                                                                                                    2'-0-Methoxyethyl (2'-MOE) nucleotides"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      nucleotides"
                          Gaps
                          ..
                                                                                                                                                                                                                                        gene therapy, antisense technology, PCTAIRE protein kinase 2, neurological disorder, human, PCTAIRE protein kinase 2; ss.
                                                                                                                                                                                                               Human PCTAIRE protein kinase 2 antisense oligonucleotide #61.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 13.8; DB 1; Length 20; Pred. No. 1e+03;
Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /note= "OTHER= 2'-O-Methoxyethyl (2'-MOE)
                          Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 20 BP; 3 A; 6 C; 3 G; 8 T; 0 U; 0 Other;
   DB 1;
   Score 13.8; DB 1
Pred. No. 1e+03;
                            Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example 15; SEQ ID NO 74; 58pp; English.
                                                                                                                                                                                                                                                                                                                                                          /note= "OTHER= Phospho:
are 5-methylcytidines"
                                                                                                                                                                                                                                                                                                        Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                 base= OTHER
                                                                                                                                                                                                                                                                                                                                                OTHER
                                                                                                                                                                                                                                                                                                                                                                                                           OTHER
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/note= "OTHER= 2
15...20
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                                                     888
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                                                                                                                                         BP
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   0.8%;
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                                                                              4 ACCTGTATGTCTGTGGG
                                                     872 ACCTGGATGACTGTGGG
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                              Conservative
                                                                                                                                            ADI19207 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                /*tag=
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                                                                                                                                                                                                                                                                                                                                                                                                                                                             /mod
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              Local Similarity
nes 15; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   US2003225256-A1
                                                                                                                                                                                                                                                                                                                       modified_base
                                                                                                                                                                                                                                                                                                                                                                                    modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                     modified base
                                                                                                                                                                                                                                                                                   Homo sapiens
                                                                                                                                                                                           22-APR-2004
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
                                                                                                                                                                    ADI19207;
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     Query Match
                                                                                                                  RESULT 1606
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                               Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /mod_base= OTHER
/note= "OTHER= Phosphorothioate backbone. All cytidines
are 5-methylcytidines"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New antisense oligonucleotide, having a sequence targeted to a nucleic acid encoding PCTAIRE protein kinase 2, useful for preparing a composition for treating neurological disorders.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /mod_base= OTHER
/note= "OTHER= 2'-0-Methoxyethyl (2'-MOE) nucleotides"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              "mod_base= OTHER
note= "OTHER= 2'-O-Methoxyethyl (2'-MOE) nucleotides"
          Gaps
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0
                                                                                                                                                                                                                                                                                            Human PCTAIRE protein kinase 2 antisense oligonucleotide #123.
                                                                                                                                                                                                                                                                                                                                   gene therapy, antisense technology, PCTAIRE protein kinase 2, neurological disorder, human, PCTAIRE protein kinase 2; ss.
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Output

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          Indels
          5
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      Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example 15; SEQ ID NO 136; 58pp; English
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      °,
                                        1436 AGGATGCCATGAAACAT 1452
                                                                                                                                                                                 BP.
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                                                                             20 AAGAGGCCATGAACAT
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                                                                                                                                                                                                                                                         (first entry)
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    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     15. .20
                                                                                                                                                                                 DNA;
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                                                                                                                                                                           ADI19269 standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                                     Key
modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       modified base
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                                                                                                                                                                                                                                                                                                                                                                                                Homo sapiens
                                                                                                                                                                                                                                                         22-APR-2004
    15;
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                                                                                                                                                                                                                    ADI19269;
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                                                                                                                                      RESULT 1607
Matches
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The invention describes a new antisense oligonucleotide, having a sequence comprising 8-80 bp targeted to a nucleic acid encoding PCTAIRE protein Kinase 2, that specifically hybridises with the nucleic acid encoding PCTAIRE protein Kinase 2 and having a sequence comprising 20 bp. The antisense oligonucleotide is useful for preparing a composition for treating e.g., neurological disorders. This sequence represents a human PCTAIRE protein kinase 2 antisense oligonucleotide.
                                                                                                                                                                                                                                                                                                                                                               /mod_base= OTHER
/note= "OTHER= Phosphorothioate backbone. All cytidines
are 5-methylcytidines"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New antisense oligonucleotide, having a sequence targeted to a nucleic acid encoding PCTAIRE protein kinase 2, useful for preparing a composition for treating neurological disorders.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               nucleotides'
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0
                                                                                                                                    Human PCTAIRE protein kinase 2 antisense oligonucleotide #67.
                                                                                                                                                                                gene therapy; antisense technology; PCTAIRE protein kinase neurological disorder; human; PCTAIRE protein kinase 2; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 13.8; DB 1; Length 20;
Pred. No. 1e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                           /*tag= a
/mcd_base= OTHER
/note= "OTHER= 2'-O-Methoxyethyl (2'-MOE)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /*tag= c
/mod_base= OTHER
/note= "OTHER= 2'-0-Methoxyethyl (2'-MOE)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  7
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                                                                                                                                                                                                                                                                                                  Location/Qualifiers
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       HP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   31-MAY-2002; 2002US-00160787.
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ADI19213 standard; DNA; 20
                                                                                         (first entry)
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                                                                                                                                                                                                                                                                                                                                        /*tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PHARM INC
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                                                                                                                                                                                                                                                                                             Key
modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                    modified_base
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                                                                                       22-APR-2004
                                                                                                                                                                                                                                                   Homo sapiens
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                                          ADI19213
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Matches
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Gaps

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1525 ATTCAGCTACAAAAGGA 1541

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antiasthmatic; respiratory; gene therapy; asthma; bronchial hyperresponsiveness; atopy; chronic obstructive lung disease; adult respiratory distress syndrome; obesity; inflammatory bowel disease; human; gene 216; single nucleotide polymorphism; SNP; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gene 216 nucleic acid, useful for preparing a composition for treating disorders e.g., asthma, bronchial hyperresponsiveness, atopy, chronic obstructive lung disease and adult respiratory distress syndrome.
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Eerdewegh PV, Dupuis J,
                                                                                                 Human gene 216 SNP detection primer seq id 94.
                       ADJ36703 standard; DNA; 20 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Pandit S;
                                                                                                                                                                                                                                                                                                        13-APR-2001; 2001US-00834597.
19-APR-2002; 2002US-00126022.
                                                                                                                                                                                                                                                                    17-OCT-2002; 2002US-00277216
                                                                                                                                                                                                                                                                                            13-APR-2000; 2000US-00548797
                                                                             22-APR-2004 (first entry)
                                                                                                                                                                                                                                                                                                                                              KEITH T.
LITTLE R D.
VAN EERDEWEGH P.
                                                                                                                                                                                                                                                                                                                                                                                               DEL MASTRO R
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Little RD,
Allen K, I
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2004-061675/06
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                           SIMON J.
ALLEN K.
                                                                                                                                                                                                                                                                                                                                                                                      DUPUIS J
                                                                                                                                                                                                                                                                                                                                                                                                                                     PANDIT S
                                                                                                                                                                                                                    US2004002470-A1.
                                                                                                                                                                                               Homo sapiens.
                                                                                                                                                                                                                                              01-JAN-2004.
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Keith T,
Simon J,
                                                      ADJ36703;
                                                                                                                                                                                                                                                                                                                                                                                               (DMAS/)
(SIMO/)
(ALLE/)
(PAND/)
                                                                                                                                                                                                                                                                                                                                                 (KEIT/)
                                                                                                                                                                                                                                                                                                                                                                       (VEER/)
                                                                                                                                                                                                                                                                                                                                                                                      ( /nand)
     RESULT 1610
                   ADJ36703
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The present invention relates to antisense compounds, compositions and methods used for modulating the expression of haem oxygenase (HO) 1. The methods and compositions of the present invention are useful for the diagnosis, prevention and/or treatment of diseases or conditions associated with aberrant expression or activity of haem oxygenase I such as hyperbilirubinaemia, neonatal jaundice and neurodegenerative diseases like Alzheimer's and Parkinson's disease. The invention is also useful in
                                                                                                                                                                                                                                                   "mod base= OTHER _{\mbox{\scriptsize Table}} " phosphorothioate backbone where all cytidines are
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           antisense-therapy. The present sequence is human haem oxygenase 1 antisense oligonucleotide used in the exemplification of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             heme
                                                                                                      Haem oxygenase 1; HO; hyperbilirubinaemia; neonatal jaundice; neurodegenerative disease; Alzheimer's disease; Parkinson's disease; antisense-therapy; nootropic; neuroprotective; human; phosphorothioate backbone; antisense; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New antisense oligonucleotide compounds, useful for diagnosing, preventing and/or treating conditions with aberrant activity of oxygenase 1, such as hyperbilirubinemia, neonatal jaundice and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ..
0
                                                                                 Human haem oxygenase 1 antisense oligonucleotide, ISIS #203104.
                                                                                                                                                                                                                                                                                                                                                                           not\overline{e}= "2'-methoxyethyl (2'-MOE) nucleotides"
                                                                                                                                                                                                                                                                                                   *tag= a
/mod_base= OTHER
/note= "2'-methoxyethyl (2'-MOE) nucleotides"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0.8%; Score 13.8; DB 1; Length 20; 88.2%; Pred. No. 1e+03; cive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 20 BP; 3 A; 7 C; 7 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example 15; SEQ ID NO 15; 43pp; English.
                                                                                                                                                                                                              Location/Qualifiers
                                                                                                                                                                                                                                                                            5-methylcytidines"
                                                                                                                                                                                                                                                                                                                                                         *tag= c
mod base= OTHER
           ADJ31669 standard; DNA; 20 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                         20-JUN-2002; 2002US-00178258
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 20-JUN-2002; 2002US-00178258
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          neurodegenerative diseases.
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Matches 15; Conservative
                                                               (first entry)
                                                                                                                                                                                                                                                                                                                                          16. .20
                                                                                                                                                                                                                                                                                            ..5
*tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                              . 20
                                                                                                                                                                                                                                         *tag=
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modified_base
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                                                             22-APR-2004
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                                                                                                                                                                                            Synthetic.
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                                      ADJ31669;
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Del Mastro RG;

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The invention describes a new isolated nucleic acid comprising a fully defined sequence having 23574 bp or at least its 50 or 15 contiguous cucleotides and includes: allele G of single nucleotide polymorphism or (SNP) AB+2; allele G of SNP BG-1; and allele G of SNP BG-2. The invention describes identifying increased susceptibility to a disorder comprising asthma, bronchial hyperresponsiveness, atopy, chronic obstructive lung disease and adult respiratory distress syndrome in a subject comprising testing a biological sample obtained from a subject for the presence of treating a biological sample obtained from a subject for the presence of the presence of the increased susceptibility to the disorder. The cucleic acid is useful for preparing a composition for treating disorders comprising asthma, bronchial hyperresponsiveness, atopy, chronic cobstructive lung disease and adult respiratory distress syndrome. This sequence represents a primer used to detect single nucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               .
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0.8%; Score 13.8; DB 1; Length 20;
88.2%; Pred. No. 1e+03;
tive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 20 BP; 5 A; 7 C; 4 G; 4 T; 0 U; 0 Other;
Example 10; SEQ ID NO 94; 441pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             polymorphisms in the human gene 216.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                554
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                538 CCCATCTTTGACAAGCC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2 cccrrcrgrgacaagcc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Best Loca
Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       à
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CCTGAAGCAGTACCTGG 877

861

à qq

20 ccregaccacaccice 4

human; single nucleotide polymorphism; SNP; ss; primer.

Synthetic.

Primer of the invention #2696.

(first entry)

06-MAY-2004

ADK96976;

BP

ADK96976 standard; DNA; 20

RESULT 16 ADK96976/

19 CCCTTCTGTGACAAGCC 3

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The invention describes a new isolated nucleic acid comprising a fully defined sequence having 23574 bp or at least its 50 or 15 contiguous nucleotides and includes: allele G of single nucleotide polymorphism (SNP) ABE; allele G of SNP BC+1; and allele C of SNP BC+2. The invention describes identifying increased susceptibility to a disorder comprising at the properties of sathma, bronchial hyperresponsiveness, atopy, chronic obstructive lung disease and adult respiratory distress syndrome in a subject comprising testing a biological sample obtained from a subject for the presence of at least one allele or haplotype given in the specification, where the nucleic acid is useful for preparing a composition for treating disorders comprising asthma, bronchial hyperresponsiveness, acopy, chronic obstructive lung disease and adult respiratory distress syndrome. This
                                                                                                                                                           antiasthmatic, respiratory; gene therapy; asthma; bronchial hyperresponsiveness; atopy; chronic obstructive lung disease; adult respiratory distress syndrome; obesity; inflammatory bowel disease; human; gene 216; single nucleotide polymorphism; SNP; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gene 216 nucleic acid, useful for preparing a composition for treating disorders e.g., asthma, bronchial hyperresponsiveness, atopy, chronic obstructive lung disease and adult respiratory distress syndrome.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Del Mastro RG;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0.8%; Score 13.8; DB 1; Length 20; 18.2%; Pred. No. 1e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 20 BP; 4 A; 4 C; 7 G; 5 T; 0 U; 0 Other;
                                                                                                                             Human gene 216 SNP detection primer seq id 154.

    Eerdewegh PV, Dupuis J,
Pandit S;

                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example 10; SEQ ID NO 154; 441pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    sequence represents a primer used to polymorphisms in the human gene 216.
                                ADJ36763 standard; DNA; 20 BP.
                                                                                                                                                                                                                                                                                                                                                                        13-APR-2000; 2000US-00548797.
13-APR-2001; 2001US-00834597.
19-APR-2002; 2002US-00126022.
                                                                                                                                                                                                                                                                                                                                           17-OCT-2002; 2002US-00277216.
                                                                                              (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match 0.89
Best Local Similarity 88.29
Matches 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                        KEITH T.
LITTLE R D.
VAN EERDEWEGH P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DUPUIS J.
DEL MASTRO R G.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Little RD,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2004-061675/06.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Allen K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SIMON J.
ALLEN K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PANDIT S
                                                                                                                                                                                                                                                                            US2004002470-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DUPUIS
                                                                                                                                                                                                                                            Homo sapiens.
                                                                                              22-APR-2004
                                                                                                                                                                                                                                                                                                            01-JAN-2004.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Keith T,
Simon J,
                                                                ADJ36763;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (SIMO/)
(ALLE/)
(PAND/)
                                                                                                                                                                                                                                                                                                                                                                                                                                        (KEIT/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (VEER/)
RESULT 1611
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                ADJ36763/c
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The present invention relates to a polynucleotide isolated from a human gene and is useful for detecting a single nucleotide polymorphism in a human gene or for diagnosing of disease. The invention enables the detection of a single nucleotide polymorphism in a human gene. The
                                                                                                                                                                                                                                                                      Novel polynucleotide useful for PCR amplification along with two DNA fragment from another set of sequences, or for detecting single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ·.
                                                                                                                                                                                                                                                                                                                                                                                                                                                    0.8%; Score 13.8; DB 1; Length 20; 88.2%; Pred. No. 1e+03; tive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         human; single nucleotide polymorphism; SNP; ss; primer.
                                                                                                                                                                                                                                                                                                                                                                                                       present sequence represents a primer of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 20 BP; 3 A; 8 C; 4 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                         Claim 2; SEQ ID NO 6005; 2627pp; Japanese.
                                                                                                                                                                                                                          (KAGA-) KAGAKU GIJUTSU SHINKO JIGYODAN.
                                                                                                                                                                                                                                                                                    fragment from another set of sequences nucleotide polymorphism in human gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        181
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           BP
                                                                                                                                                                          08-MAR-2002; 2002JP-00064373.
                                                                                                                                                                                                 08-MAR-2002; 2002JP-00064373
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Primer of the invention #523.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              19 AGTCCGAGGTGGCCAAG 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              08-MAR-2002; 2002JP-00064373.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      08-MAR-2002; 2002JP-00064373
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    165 ACTCCGAGGTGGCCGAG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADK94803 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          06-MAY-2004 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match 0.8
Best Local Similarity 88.2
Matches 15; Conservative
                                                                                                                                                                                                                                                   WPI; 2004-093977/10.
                                                                                                                        JP2003259875-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            JP2003259875-A.
                                                                                                                                                 16-SEP-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Synthetic
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Gaps

0;

Indels

2;

0; Mismatches

88.2%;

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BP; 3 A; 8 C; 4 G; 5 T; 0 U; 0 Other,
                                                                                                                                                                                                                                                          Oligonucleotide associated to IL5R-X61176 #85.
                                                             Claim 2; SEQ ID NO 3832; 2627pp; Japanese
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 2; SEQ ID NO 2249; 85pp; English.
    (KAGA-) KAGAKU GIJUTSU SHINKO JIGYODAN
                                               nucleotide polymorphism in human gene.
                                                                                                                                                              1729 CACCTGCCCACTTGTCC 1745
                                                                                                                                                                            1 cacgreaccacrrerce 17
                                                                                                                                                                                                                                                                                                                                                            25-JUL-2003; 2003WO-US023509
                                                                                                                                                                                                                                                                                                                                                                          29-JUL-2002; 2002US-0399076P
                                                                                                                                                                                                                                                                                                                                                                                        (EPIG-) EPIGENESIS PHARM INC
                                                                                                                                                                                                                ADJ61393 standard; DNA; 20
                                                                                                                                         Local Similarity 88.2
Les 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                disease e.g., asthma.
                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2004-203534/19.
                   WPI; 2004-093977/10
                                                                                                                                                                                                                                                                                                                                 WO2004011613-A2
                                                                                                                                                                                                                                                                                                                                                                                                      Nyce JW, Tang
Shahabuddin S,
                                                                                                                                                                                                                                                                                                                    Homo sapiens.
                                                                                                                                                                                                                                                                                                                                               05-FEB-2004
                                                                                                                      Sequence 20
                                                                                                                                                                                                                                              06-MAY-2004
                                                                                                                                   Query Match
                                                                                                                                                                                                                               ADJ61393
                                                                                                                                                                                                  RESULT 1614
ADJ61393
                                                                                                                                            Best Loca
Matches
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à
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(first entry)

BP.

ed. No. 1e+03; Mismatches

0

Score 13.8; Pred. No. 1e

0.8%;

Aguilar D,

L, Sandrasagra A, Lu H, Cong H;

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The present invention relates to an oligonucleotide anti-sense to e.g., initiation codon, coding region with 2-10 nucleotides of 5'-end and 3'-end of nucleic acid target comprising gene(s) chosen from e.g. interleukin (IL)-4 receptor, IL-5 receptor or salts of the oligonucleotide and optionally surfactant operatively linked to the oligonucleotide. The method is useful for preventing or treating a respiratory or lung disease, which involves administering to the airways of a subject an effective amount of an inhibitor. The oligonucleotide is useful for production of a medicament for the prevention and/or treatment of a respiratory or lung disease. The respiratory or lung disease is
            respiratory or lung disease, which involves administering to the dirways of a subject an effective amount of an inhibitor. The oligonuclectide is useful for production of a medicament for the prevention and/or treatment of a respiratory or lung disease. The respiratory or lung disease. The respiratory or lung disease. The respiratory or lung disease is chosen from alrway inflammation, allergy(les), asthma, impeded respirator, cystic fibrosis (CP), chronic obstructive pulmonary diseases (COPD), allergic thinitis (AR), acute respiratory distress syndrome (ARDS), pulmonary hypertension, lung inflammation, bronchitis, alrway obstruction. The present sequence represents an oligonucleotide of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             e.g.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  interleukin; IL-4 receptor; IL-5 receptor; lung disease; airway inflammation; allergy; asthma; impeded respiration; cystic fibrosis; acute respiratory distress syndrome; pulmonary hypertension; lung inflammation; bronchitis; oligonucleotide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Novel single or multiple target oligonucleotide anti-sense to e.g. initiation codons and introns of respiratory disease-relevant genes cCRI, RANTES, MCP4, useful for prophylaxis or treating respiratory disease e.g., asthma.
                                                                                                                                                                                                                                                                                                                                                                               Gaps
oligonucleotide. The method is useful for preventing or treating a
                                                                                                                                                                                                                                                                                                                                                                               .
                                                                                                                                                                                                                                                                                                                                Length 20;
                                                                                                                                                                                                                                                                                                                                                                               Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ŝ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Miller
                                                                                                                                                                                                                                                                                    Sequence 20 BP; 6 A; 4 C; 3 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                0.8%; Score 13.8; DB 1;
88.2%; Pred. No. 1e+03;
                                                                                                                                                                                                                                                                                                                                                                               2
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                                                                                                                                                                                                                                                                                                                                                      Pred. No. 1e+03;
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Oligonucleotide associated to IL 5R #149.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             claim 2; SEQ ID NO 308; 85pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sandrasagra A,
H, Cong H;
                                                                                                                                                                                                                                                                                                                                                                                                                                   1444 ATGAAACATCCATTCTT 1460
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (EPIG-) EPIGENESIS PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           25-JUL-2003; 2003WO-US023509.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          29-JUL-2002; 2002US-0399076P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   70
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Argaagcarccaractr
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DNA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Nyce JW, Tang L, San
Shahabuddin S, Lu H,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2004-203534/19.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADJ59452 standard;
                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity
Matches 15; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WO2004011613-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   06-MAY-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   05-FEB-2004
                                                                                                                                                                                                                                                    invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADJ59452;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RESULT 1615
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADJ5945
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            8×3000000000×8
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0
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initiation codons and introns of respiratory disease-relevant genes e.g.,
CCR1, RANTES, MCP4, useful for prophylaxis or treating respiratory
                                                                                                                                                                                                                                                                     The present invention relates to a polynucleotide isolated from a human gene and is useful for detecting a single nucleotide polymorphism in a human gene or for diagnosing of disease. The invention enables the detection of a single nucleotide polymorphism in a human gene. The present sequence represents a primer of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  interleukin, IL-4 receptor, IL-5 receptor; lung disease, airway inflammation; allergy; asthma; impeded respiration; cystic fibrosis; acute respiratory distress syndrome; pulmonary hypertension; lung inflammation; bronchitis; oligonucleotide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The present invention relates to an oligonucleotide anti-sense to eninitiation codon, coding region with 2-10 nucleotides of 5'-end and end of nucleic acid target comprising gene (8) chosen from e.g. interleukin (IL)-4 receptor, IL-5 receptor or salts of the oligonucleotide and optionally surfactant operatively linked to the
                                                                                                                                two DNA
                                                                                                                                Novel polynucleotide useful for PCR amplification along with two fragment from another set of sequences, or for detecting single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ..
0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DB 1; Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Indels
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pulmonary diseases

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Gaps

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The invention relates to an insect acetylcholine esterase (ace-1) (1) with a central catalytic region, given in the specification as AbJ93656, or (ii) a sequence 60 % identical or 70 % similar to AbJ93656. Also disclosed is a method for detecting insects that carry resistance to organophosphorus and carbamate insecticides. The method of the invention sessitant to organophosphorus and catbamate insecticides. Ace-1 and transgenic invertebrates that express it, are used to screen for insecticides, i.e. inhibitors of ace-1. The nucleic acid that encodes (1), also antibodies specific for (1), are used to detect insects, that are resistant to organophosphorus and carbamate insecticides. Secondary continuation of ace-1. The nucleic acid that encodes particularly mosquito disease vectors but also agricultural pests, that are resistant to organophosphorus and carbamate insecticides. Agents that are resistant to organophosphorus and carbamate insecticides. Agents that
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         inhibit (I) are effective against insect strains resistant to organophosphorus and carbamate insecticides. Sequences given in ADJ93657-ADJ93784 represent products of the ace-1 gene (CDNA, protein, AChE1) from various insects, and also primers for the amplification of ace-1 nucleic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Insecticide; acetylcholine esterase; ace-1; organophosphorus; carbamate;
insecticide; resistance; AChEl; mosquito; PCR; primer; ss.
chosen from airway inflammation, allergy(ies), asthma, impeded respiration, cystic fibrosis (CP), chronic obstructive pulmonary diseas (COPD), allergic rhinitis (AR), acute respiratory distress syndrome (ARDS), pulmonary hypertenalon, lung inflammation, bronchitis, airway obstruction. The present sequence represents an oligonucleotide of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New insect acetylcholine esterase, useful in screening for insecticides effective against strains resistant to organophosphates and carbamates.
                                                                                                                                                                         Score 13.8; DB 1; Length 20;
Pred. No. 1e+03;
0; Mismatches 2; Indels
                                                                                                                                 Sequence 20 BP; 6 A; 4 C; 4 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                              0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Pasteur N;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 11; SEQ ID NO 123; 169pp; French.
                                                                                                                                                                                                                                                      1444 ATGAAACATCCATTCTT 1460
                                                                                                                                                                                                                                                                              3 ATGAAGCATCCATACTT 19
                                                                                                                                                                                                                                                                                                                                                                                          BP.
                                                                                                                                                                           0.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Fort P, Raymond M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2003WO-FR001876.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2002FR-00007622.
2002FR-00013799.
                                                                                                                                                                                                                                                                                                                                                                                        20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (CNRS ) CENT NAT RECH SCI. (UYMO-) UNIV MONTPELLIER 2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
                                                                                                                                                                                                              15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                    ADJ93778 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AG
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                                                                                                                                                                                             Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Forward primer Ex3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WO2004000994-A2,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                19-JUN-2003;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                 06-MAY-2004
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                                                                                                 invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                           ADJ93778;
                                                                                                                                                                         Query Match
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Sequence 20 BP; 3 A; 5 C; 7 G; 5 T; 0 U; 0 Other;

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New antisense compounds targeted to nucleic acid molecules encoding phospholipase D2, useful for treating diseases associated with expression of phospholipase D2, e.g. cancer, Alzheimer's disease or Parkinson's
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The present invention relates to antisense oligonucleotides which are targetted to nucleic acid molecule encoding phospholipase D2 and the encoding protein. The invention is useful for inhibiting the expression of phospholipase D2 and for treating diseases and conditions associated with expression of phospholipase D2 e.g. hyperproliferative disease such as Alzheimer's disease and Parkinson's disease. The invention is also useful for therapeutic and prophylactic applications to prevent or delay infection, inflammation and tumour formation. The present sequence is human phospholipase D2 target
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ;
0
                                     Gaps
                                                                                                                                                                                                                                                                                      Phospholipase D2; hyperproliferative disorder; cancer;
neurodegenerative disease; Alzheimer's disease; Parkinson's disease;
infection; inflammation; tumour; therapy; human; ss.
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   Score 13.8; DB 1; Length 20;
Pred. No. 1e+03;
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88.2%; Pred. No. 1e+03;
ive 0; Mismatches 2; Indels
                                  Indels
                                 5;
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                                                                                                                                                                                                                                                         Human phospholipase D2 target oligonucleotide
                               0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example 15; SEQ ID NO 51; 46pp; English.
                                                              1611
                                                                                   3 TCGTGGACACCGTGTTC 19
                                                                                                                                                                      BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADJ64112 standard; DNA; 20 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                          20-JUN-2002; 2002US-00177896.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         20-JUN-2002; 2002US-00177896.
   0.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 88.2%;
                                                                                                                                                                 ADJ64147 standard; DNA; 20
                                                           1595 TGGTGGACACCGAGTTC
                                                                                                                                                                                                                             (first entry)
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Query Match

Best Local Similarity 88.2
Matches 15; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Dobie KW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2004-081729/08.
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                                                                                                                                                                                                                                                                                                                                                     Homo sapiens.
                                                                                                                                                                                                                             06-MAY-2004
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                                                                                                                                    RESULT 1617
ADJ64147/c
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human; ss; liver related homologue-1; LRH1; NR5A2; antisense; phosphorothioate; 2' MOE, breast cancer; dyslipidaemia; atherosclerosis; low HDL; high density lipoprotein; high LDL; hypercholesterolaemia; gall stone; trigityceridaemia; obesity; heptitis; heptitis; heptitocellular carcinoma; arcomatase; cytostatic; antilipaemic; antiarteriosclerotic; anorectic; hepatotropic; litholytic; antiinflammatory; virucidal.

sapiens

Homo

Synthetic.

Antisense DNA oligo used to modulate human LRH1 expression SeqID 335.

(first entry)

20-MAY-2004

ADJ15785;

ADJ15785 standard; DNA; 20 BP.

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The present invention relates to antisense oligonucleotides which are targetted to nucleic acid molecule encoding phospholipase D2 and the encoding protein. The invention is useful for inhibiting the expression of phospholipase D2 and for treating diseases and conditions associated with expression of phospholipase D2 e.g. hyperproliferative disorder such as cancer, neurodegenerative disease such as Alzheimer's disease and parkingon's disease. The invention is also useful for therapeutic and prophylactic applications to prevent or delay infection, inflammation and tumour formation. The present sequence is human phospholipase D2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New antisense compounds targeted to nucleic acid molecules encoding phospholipase D2, useful for treating diseases associated with expression of phospholipase D2, e.g. cancer, Alzheimer's disease or Parkinson's
                                                                                                                                                                                                                                    not\,\bar{e}\text{=} "Phosphorothioate backbone where all cytidines are 5'-methylcytidines"
                                                                               Phospholipase D2; hyperproliferative disorder; cancer;
neurodegenerative disease; Alzheimer's disease; Parkinson's disease;
                                                                                                          infection; inflammation; tumour; therapy; human; antisense; ss.
                                                                                                                                                                                                                                                                                                                  "2'- methoxyethyl (2'-MOE) nucleotides"
                                                  Human phospholipase D2 antisense oligonucleotide ISIS #159040.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 13.8; DB 1; Length 20;
Pred. No. 1e+03;
0; Mismatches 2; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 20 BP; 5 A; 3 C; 9 G; 3 T; 0 U; 0 Other;
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                                                                                                                                                                                   Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                              /mod_base= OTHER
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                           (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                              US2004005705-A1
                                                                                                                                                                                                                                                                                                                                      modified base
                                                                                                                                                                                                      modified base
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                                                                                                                                            sapiens
                           06-MAY-2004
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                                                                                                                                                                                                                                                                                                                                                                                                                                         08-JAN-2004
                                                                                                                                                             Synthetic.
                                                                                                                                               Homo
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nucleotides"

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This invention relates to novel antisense compounds useful for modulating the expression of liver related homologue-1 (LRH1) and splice variants thereof. Specifically, it refers to compositions 8-30 nucleobases in length that target a portion of an active site on the nucleic acid conclear receptor protein that functions as a tissue specific conclear receptor protein that functions as a tissue specific cranscription factor. The present invention describes antisense of inkage, a phosphorothiate linkage, at least one modified internucleoside oligonucleotides that comprise at least one modified sugar moiety, as 2'-0-methoxyethyl (2' MOE) and at least one modified sugar moiety, concept, as a speciated with LRH1, such as breast cancer, diagnosing a disease associated with LRH1, such as breast cancer, diagnosing a disease associated with LRH1, such as breast cancer, diagnosing a disease associated with LRH1, such as breast cancer, cut ilphyceridaemia, obesity, hepatitis B virus-mediated acute or chronic triglyceridaemia, obesity, hepatitis B virus-mediated acute or chronic hepatitis, as well as hepatocellular carcinoma or a condition associated with aromatase activity. Accordingly, these compositions exhibit cycleatic, antilipaemic, antilarterlosclerotic, anorectic, hepatorropic,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New antisense oligonucleotides targeted to a nucleic acid encoding liver related homologue-1 (LRH1), useful for treating breast cancer, dyslipidemia, atherosclerosis, hypercholesterolemia, or hepatitis.
                                                                                                                                                                                                                                                                                                                                                                                                                                          /note= "OTHER= 2' methoxyethyl (2' MOE) nucleotides. All cytidine nucleobases are 5-methylcytidine."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /note= "OTHER= 2' methoxyethyl (2' MOE) nucleotides. All
cytidine nucleobases are 5-methylcytidine."
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/label= OTHER= phosphorothioate backbone
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/note= "OTHER= 2'
                                                                                                                                                                                                                                                                                                                                                              OTHER
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/note= "OTHER= 2'
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                                                                                                                                                                                                                                                                                                                   Key
modified_base
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Gaps

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0;

Conservative

15;

Best Loc Matches

210 GCAGATAGGCCTGGATG 226

11

||||| |GCAGAATGGCCTGGATG

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RESULT 1619

ADJ15785/c

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08-JAN-2004
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  This invention relates to novel antisense compounds useful for modulating
                                                                                                                                                                                                                                   human; ss; liver related homologue-1; LRH1; NR5A2; antisense; phosphorothioate; 2' MOE; breast cancer; dyslipidaemia; atherosclerosis; low HDL; high density lipoprotein; high LDL; hypercholesterolaemia; gall stone; triglyceridaemia; obesity; hepatitis; hepatocellular carcinoma; aromatase; cytostatic; antilipaemic; antiarteriosclerotic; anorectic; hepatocropic; litholytic; antiinflammatory; virucidal.
                                                                                                                                                                                                                                                                                                                                                                                                                    /note= "OTHER= 2' methoxyethyl (2' MOE) nucleotides. All cytidine nucleobases are 5-methylcytidine."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New antisense oligonucleotides targeted to a nucleic acid encoding liver related homologue-1 (LRH1), useful for treating breast cancer, dyslipidemia, atherosclerosis, hypercholesterolemia, or hepatitis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /note= "OTHER= 2' methoxyethyl (2' MOE) nucleotides. All
litholytic, antiinflammatory and virucidal activities. This oligonucleotide sequence is an antisense DNA oligo used to modulate the expression of the human LRH1 protein of the invention.
                                                                                                                                                                                                                  Antisense DNA oligo used to modulate human LRH1 expression SeqID 3068.
                                                                               Gaps
                                                                              0;
                                                           Length 20;
                                                   Score 13.8; DB 1; Dengar-
Pred. No. 1e+03; 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         cytidine nucleobases are 5-methylcytidine.
                                                                                                                                                                                                                                                                                                                                                                               'label = OTHER = phosphorothioate backbone
                                      Sequence 20 BP; 1 A; 7 C; 5 G; 7 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                            location/Qualifiers
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                                                                                              869
                                                                                                                                                               BP.
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                                                                  88.2%;
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                                                                                              853 GACAAGGACCTGAAGCA
                                                                                                          17 GACAGGGCCCTGAAGCA
                                                                                                                                                     (first entry)
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                                                                            Conservative
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                                                                   Local Similarity
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                                                         Query Match
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                                                                                                                                     Kane CD;
                                                                          datches
5555X88
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the expression of liver related homologue-1 (LRH1) and splice variants thereof. Specifically, it refers to compositions 8-30 mucleobases in length that target a portion of an active site on the nucleic acid molecule encoding LRH1 (also known as NR5A2). LRH1 is a monomeric orphan nuclear receptor protein that functions as a tissue specific transcription factor. The present invention describes antisense oligonucleotides that comprise at least one modified internucleoside linkage, a phosphorothicate linkage, at least one modified internucleoside linkage, a phosphorothicate linkage; at least one modified sugar moiety, a 2'-O-methoxyethyl (2' MOE) and at least one modified sugar moiety, a 2'-O-methoxyethyl (2' MOE) and at least one modified sugar moiety, complycitation. These antisense compounds are useful for treating or diagnosing a disease associated with LRH1, such as breast cancer, diagnosing a disease associated with LRH1, such as breast cancer, diagnosing a disease associated with LRH1 such as breast cancer, complycationary lipoprotein), hypercholesterolaemia, gall stones, triglyceridaemia, obesity, hepatitis B virus-mediated acute or chronic hepatitis, as well as hepatocellular carcinoma or a condition associated critical antilipaemic, antilatratiosclerotic, ancreatic, antilitammatory and virucidal activities. This correctic, hepatotropic, clitholytic, antilinfammatory and virucidal activities. This colligonucleotide sequence is an antisense DNA oligo used to modulate the expression of the human LRH1 protein of the invention.
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ses are 5-methylcytidine."
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Antisense DNA oligo used to modulate human LRH1 expression SeqID 3349.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0.8%; Score 13.8; DB 1; Length 20; 88.2%; Pred. No. 1e+03; ive 0; Mismatches 2; Indels
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/note= "OTHER= 2' methoxyethyl (2' MOE) nu
cytidine nucleobases are 5-methylcytidine.
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modified_base
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADJ18799;
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nucleotides. All

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This invention relates to novel antisense compounds useful for modulating the expression of liver related homologue-1 (LRH1) and splice variants thereof. Specifically, it refers to compositions 8-30 nucleobases in compensive that target a portion of an active site on the nucleic acid composured to the nuclear receptor protein that functions as a tissue specific conclear receptor protein that functions as a tissue specific conclear receptor protein that functions as a tissue specific composition actor. The present invention describes antisense composition are useful for treatured of linkage, a phosphorothioate linkage; at least one modified internucleoside concept a 2-7-0-methoxycythyl (2' NOE) and at least one modified sugar moiety, an expression of the seast cancer, and agnosing a disease associated with LRH1, such as breast cancer, concept and adjacental actions and condition associated activity, accordingly, these compositions exhibit contained activity, Accordingly, these compositions exhibit contained and an analysis and all and an analysis and virucidal activities of the contained activity and virucidal activities. This
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New antisense oligonucleotides targeted to a nucleic acid encoding liver related homologue-1 (LRH1), useful for treating breast cancer, dyslipidemia, atherosclerosis, hypercholesterolemia, or hepatitis.
                                                                                                                                                                                                                                      /note= "OTHER= 2' methoxyethyl (2' MOE) nucleotides. All cytidine nucleobases are 5-methylcytidine."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            oligonucleotide sequence is an antisense DNA oligo used to modulate the
                                                                                                                  /note = "OTHER = 2' methoxyethyl (2' MOE) nu
cytidine nucleobases are 5-methylcytidine.
                    'label = OTHER = phosphorothioate backbone
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0.8%; Score 13.8; DB 1
38.2%; Pred. No. 1e+03;
ive 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example 15; SEQ ID NO 216; 909pp; English.
                                                                        /*tag= a
/mod_base= OTHER
                                                                                                                                                                                                                               /mod_base= OTHER
mod base= OTHER
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (PHAA ) PHARMACIA CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2004-083058/08.
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                                                      modified base
                                                                                                                                                                             modified base
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Kane CD;
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                                                                                                                                                                                                                                                                                                                                                                                          This invention relates to novel antisense compounds useful for modulating the expression of liver related homologue-1 (LAHI) and splice variants thereof. Specifically, it refers to compositions 8-30 nucleobases in length that target a portion of an active site on the nucleic acid molecule encoding LAHI (also known as NEAR). LAHI is a monomeric orphan nuclear receptor protein that functions as a tissue specific configuration factor. The present invention describes antisense coligonucleotides that comprise at least one modified internucleoside linkage, a phosphorothicate linkage; at least one modified sugar moiety, a 2-0-methoxyethyl (2' MOE) and at least one modified sugar moiety, configuration these antisense compounds are useful for treating or diagnosing a disease associated with LRHI, such as breast cancer, dispipidaemia, atherosclerosis, low HDL (high density lipoprotein), hypercholesterolaemia, gall stones, configurated acute or chronic configuration, as well as hepatocellular carcinoma or a condition associated with aromatese activity. Accordingly, these compositions exhibit cytostatic, antiliflammatory and virucidal activities. This importion is the invention of the modulate the configuration and antisense DNA oligo used to modulate the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ;
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                                                                                                                                                                                                                                                      New antisense oligonucleotides targeted to a nucleic acid encoding liver related homologue-1 (LRH1), useful for treating breast cancer, dyslipidemia, atherosclerosis, hypercholesterolemia, or hepatitis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ss; liver related homologue-1; LRH1; NR5A2; antisense;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0.8%; Score 13.8; DB 1; Length 20; 88.2%; Pred. No. 1e+03; 2; Indels ive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 expression of the human LRH1 protein of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 20 BP; 8 A; 5 C; 4 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                              Example 15; SEQ ID NO 3349; 909pp; English.
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          01-JUL-2003; 2003WO-US020865.
                                                              01-JUL-2002; 2002US-0392813P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                antiinflammatory; virucidal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1 AGGTCTAAGACACAGAC
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/*tag=
                                                                                                            (PHAA ) PHARMACIA CORP
                                                                                                                                                                                                               WPI; 2004-083058/08.
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Matches 15; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 modified base
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                                                                                                                                                                   Kane CD;
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ADJ15666,

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Gaps

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2; Indels

DB 1; Length 20;

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Kane CD
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phosphorothicate; 2' MOB; breast cancer; dyslipidaemia; atherosclerosis; low HDL; high density lipoprotein; high LDL; hypercholesterolaemia; gall stone; triglyceridaemia; obesity; hepatitis;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        nucleotides. All
Antisense DNA oligo used to modulate human LRH1 expression SeqID 3222.
                                                                                                                                    heparocellular carcinoma; arcmatase; cytostatic; antilipaemic; antiarteriosclerotic; anorectic; heparotropic; litholytic; antilinflammatory; virucidal.
                                             human; ss; liver related homologue-1; LRH1; NR5A2; antisense;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      'label= OTHER= phosphorothioate backbone
                                                                                                                                                                                                                                                                                                             Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                              base= OTHER
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modified_base
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                                                                                                                                                                                                                                       sapiens
                                                                                                                                                                                                                                                          Synthetic
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WO2004003201-A2

01-JUL-2003; 2003WO-US020865.

01-JUL-2002; 2002US-0392813P.

(PHAA) PHARMACIA CORP

WPI; 2004-083058/08.

New antisense oligonucleotides targeted to a nucleic acid encoding liver related homologue-1 (LRH1), useful for treating breast cancer, dyslipidemia, atherosclerosis, hypercholesterolemia, or hepatitis,

Example 15; SEQ ID NO 3222; 909pp; English

01-JUL-2003; 2003WO-US020865. 01-JUL-2002; 2002US-0392813P.

08-JAN-2004.

(PHAA) PHARMACIA CORP

WPI; 2004-083058/08

Kane CD;

This invention relates to novel antisense compounds useful for modulating the expression of liver related homologue-1 (LRH1) and splice variants thereof. Specifically, it refers to compositions 8-30 nucleobases in molecule encoding LRH1 (also known as NESAS). LRH1 is a monomeric orphan nuclear receptor protein that functions as a tissue specific transcription factor. The present invention describes antisense oligonucleotides that comprise at least one modified internucleoside linkage, a phosphorothioate linkage; at least one modified sugar moiety, a 2'-O-methoxyethy! (2' MOE) and at least one modified nucleobase, a 5-methylcytidine. These antisense compounds are useful for treating or mittylcytidine. These antisense compounds are useful for treating or diagnosing a disease associated with LRH1, such as breast cancer, disklipideamia, atherosclerosis, low HDL (high density lipoprotein), high triglyceridaemia, obesity, hepatitis B virus-mediated acute or chronic hepatitis, as well as hepatocellular carcinoma or a condition associated with aromatase activity. Accordingly, these compositions exhibit cytostatic, antilipaemic, antiarteriosclerotic, anorectic, hepatotropic, litholytic, antiinflammatory and virucidal activities. This oligonucleotide sequence is an antisense DNA oligo used to modulate the expression of the human LRH1 protein of the invention. oligonucleotide

Sequence 20 BP; 7 A; 5 C; 5 G; 3 T; 0 U; 0 Other;

This invention relates to novel antisense compounds useful for modulating the expression of liver related homologue-1 (LRH1) and splice variants thereof. Specifically, it refers to compositions 8-30 nucleobases in length that target a portion of an active site on the nucleic acid molecule encoding LRH1 (also known as NRBA2). LRH1 is a monomeric orphan nuclear receptor protein that functions as a tissue specific

New antisense oligonucleotides targeted to a nucleic acid encoding liver related homologue-1 (LRH1), useful for treating breast cancer, dyslipidemia, atherosclerosis, hypercholesterolemia, or hepatitis.

Example 15; SEQ ID NO 3393; 909pp; English.

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0;
                                                                                                                                                                                                human; ss; liver related homologue-1; LRH1; NR5A2; antisense; phosphorothioate; 2' MOE; breast cancer; dyslipidaemia; atherosclerosis; low HDL; high density lipoprotein; high LDL; hypercholesterolaemia; gall stone; triglyceridaemia; obesity; hepatitis; hepatocellular carcinoma; aromatase; cytostatic; antilipaemic; antiarteriosclerotic; anorectic; hepatotropic; litholytic; antiinflammatory; virucidal.
                                                                                                                                                                                                                                                                                                                                                                                                                                  /mod_base= OTHER /nethoxyethyl (2' MOE) nucleotides. All cytidine nucleobases are 5-methylcytidine."
                                                                                                                                                                                                                                                                                                                                                                                               nucleotides. All
                                                                                                                                                                             Antisense DNA oligo used to modulate human LRH1 expression SeqID 3393
                              Gaps
                               ..
0
         Score 13.8; DB 1; Length 20;
Pred. No. 1e+03;
0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                             /note= "OTHER= 2' methoxyethyl (2' MOE) nuc
cytidine nucleobases are 5-methylcytidine."
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/mod_base= OTHER
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                                               1607 AGTICTAAGCCACAGAC 1623
                                                                    4 AGGTCTAAGACACAGAC 20
                                                                                                                     BP.
       0.8%;
                                                                                                                  ADJ18843 standard; DNA; 20
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Query Match
Best Local Similarity
Watches 15; Conserve
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                                                                                                                                                                                                                                                                                  sapiens
                                                                                                                                                                                                                                                                                           Synthetic.
                                                                                                                                      ADJ18843;
                                                                                                RESULT 1624
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Kane CD;

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transcription factor. The present invention describes antisense oligonucleotides that comprise at least one modified internucleoside linkage, a phosphorothicate linkage; at least one modified sugar moiety, at 0.0 methoxyethyl (2' MOE) and at least one modified sugar moiety, at 0.0 methoxyethyl (2' MOE) and at least one modified nucleobase, a 5-methoxyethyl or these antisense compounds are useful for treating or diagnosing a disease associated with LRH1, such as breast cancer, tyslicity anterosciated with LRH1, such as breast cancer, LDL (low density lipoprotein), hypercholesterolaemia, gall stones, triglyceridaemia, obesity, hepatitis B virus-mediated acute or chronic hepatitis, as well as hepatocellular carcinoma or a condition associated with aromatase activity. Accordingly, these compositions exhibit cycostatic, antilipaemic, antiarteriosclerotic, anorectic, hepatotropic, litholytic, antiinflammatory and virucidal activities. This
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /note= "OTHER= 2' methoxyethyl (2' MOE) nucleotides. All cytidine nucleobases are 5-methylcytidine."
                                                                                                                                                                                                                                oligonucleotide sequence is an antisense DNA oligo used to modulate the expression of the human LRH1 protein of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Antisense DNA oligo used to modulate human LRH1 expression SeqID 303.
                                                                                                                                                                                                                                                                                                                                                             Gaps
                                                                                                                                                                                                                                                                                                                                                             0;
                                                                                                                                                                                                                                                                                                                          0.8%; Score 13.8; DB 1; Length 20;
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label= OTHER= phosphorothioate backbone
                                                                                                                                                                                                                                                                                                                                                             Indels
                                                                                                                                                                                                                                                                                                                                                             2;
                                                                                                                                                                                                                                                                                       Sequence 20 BP; 8 A; 6 C; 4 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                          Pred. No. 1e+03;
0; Mismatches
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'mod base= OTHER
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                                                                                                                                                                                                                                                                                                                                                                                                                                 AGGTCTAAGACACAGAC 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADJ15753 standard; DNA; 20 BP.
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Best Local Similarity 88.2
Matches 15; Conservative
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This invention relates to novel antisense compounds useful for modulating the expression of liver related homologue-1 (LRH1) and splice variants thereof. Specifically, it refers to compositions 8-30 nucleobases in clearth that target a portion of an active site on the nucleic acid molecule encoding LRH1 (laso known as NR5A2). LRH1 is a monomeric orphan complear receptor protein that functions as a tissue specific runlear receptor protein that functions as a tissue specific coligonucleotides that comprise at least one modified internucleoside clinkage, a phosphorothicate linkage; at least one modified internucleoside clinkage, a phosphorothicate linkage; at least one modified sugar moiety, a 2-0-methoxyethyl (2' MOE) and at least one modified nucleobase, a 5-cc methylcytidine. These antisense compounds are useful for treating or diagnosing a disease associated with LRH1, such as breast cancer, diagnosing a disease associated with LRH2, such as breast cancer, complication and a shear or she with a translated acute or chronic triglyceridemia, obesity, hepatitis B virus-mediated acute or chronic curing and an antistant arctinoma or a condition associated with armates antistic manner and antistant antistant and antistic manner.
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                                                                                    New antisense oligonuclectides targeted to a nucleic acid encoding liver related homologue-1 (LRH1), useful for treating breast cancer, dyslipidemia, atherosclerosis, hypercholesterolemia, or hepatitis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              oligonucleotide sequence is an antisense DNA oligo used to modulate the expression of the human LRH1 protein of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Antisense DNA oligo used to modulate human LRH1 expression SeqID 3427.
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litholytic, antiinflammatory and virucidal activities. This
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/*tag= b
/*nd base= OTHER
/label= OTHER= phosphorothioate backbone
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                                                                                                                                                                        Example 15; SEQ ID NO 303; 909pp; English.
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modified_base
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Synthetic.
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This invention relates to novel antisense compounds useful for modulating the expression of liver related homologue-1 (LRHI) and splice variants therefor Specifically, it refers to compositions 8-30 nucleobases in length that target a portion of an active site on the nucleic acid molecule encoding LRHI (also known as NRSA2). LRHI is a monomeric orphan conclude receptor protein that functions as a tissue specific transcription factor. The present invention describes antisense oligonucleotides that comprise at least one modified internucleoside linkage, a phosphorothioate linkage; at least one modified sugar moiety, a 2'-O-methoxyethyl (2' MOE) and at least one modified sugar moiety, a 2'-O-methoxyethyl (2' MOE) and at least one modified nucleobase, a 5-comethoxyethyl (2' MOE) and at least one modified nucleobase, a 5-comethoxyethyl (2' MOE) and at least one modified nucleobase, a 5-comethoxyethyl (2' MOE) and at least one modified nucleobase, a 5-comethoxyethyl (2' MOE) and at least one modified nucleobase, a 5-comethoxyethyl at these antisense compounds are useful for treating or diagnosing a disease associated with LRHI, such as breast cancer, dyslipidaemia, obesity, hepatitis B virus-mediated acute or chronic hepatitis, as well as hepatociallar carcinoma or a condition associated with aromatase activity. Accordingly, these compositions exhibit control of the aromatase activity. Accordingly these compositions exhibit and relations and virusidal activities of the aromatase activity. Accordingly and virusidal activities of the aromatase activity. Accordingly these compositions exhibit and a structure of the aromatase activity.
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    /note= "OTHER= 2' methoxyethyl (2' MOE) nucleotides. All
                                                                                                                       /note= "OTHER= 2' methoxyethyl (2' MOE) nucleotides. All cytidine nucleobases are 5-methylcytidine."
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                             cytidine nucleobases are 5-methylcytidine.
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88.2%; Pred. No. 1e+03;
ative 0; Mismatches
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                                                                                                    base= OTHER
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ID AD017468 standard; DNA; 20 BP.

XC AD017468;

XX AC AD017468;

XX D-MAY-2004 (first entry)

XX DE Antisense DNA oligo used to modu;

XX XX Numan; 85; liver related homology

XW human; 85; liver related homology

XW phosphorothioate; 2' MOE; breast
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Best Local Similarity 88.2°
Warches 15; Conservative
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human; ss, liver related homologue-1; LRH1; NR5A2; antisense; phosphorothicate; 2' MOE; breast cancer; dyslipidaemia; atherosclerosis;

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This invention relates to novel antisense compounds useful for modulating the expression of liver related homologue-1 (LRH1) and splice variants thereof. Specifically, it refers to compositions 8-30 nucleobases in length that target a portion of an active site on the nucleic acid molecule encoding LRH1 (also known as NESA2). LRH1 is a monomeric orphan culled receptor protein that functions as a tissue specific transcription factor. The present invention describes antisense oligonucleotides that comprise at least one modified internucleoside linkage, a phosphorothicate linkage; at least one modified nucleobase, a 2.0-methoxyethyl (2. MOE) and at least one modified nucleobase, a 5-comethoxyethyl (2. MOE) and at least one modified nucleobase, a 5-comethoxyethyl (2. MOE) and at least one modified nucleobase, a 5-comethoxyethyl at antisense compounds are useful for treating or diagnosing a disease associated with LRH1, such as breast cancer, constipulatemia, atheroscalerosis, low HDL (high density lipoprotein), hypercholesterolaemia, gall stones, triglyceridaemia, obesity, hepatitis B virus-mediated acute or chronic with argument and associated with LHH argument and associated with argument are condition associated with argument and acute or chronic with argument and associated with argument and associated with argument and a secondated.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
ow HDL; high density lipoprotein; high LDL; hypercholesterolaemia;
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                   gall stone; triglyceridaemia; obesity; hepaticis; hepaticis; hepaticis; hepaticis; antilipaemic; antiarteriosclerotic; anorectic; hepatotropic; litholytic; antiinflammatory; virucidal.
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0.8%; Score 13.8; DB 1; Length 20;
Best Local Similarity 88.2%; Pred. No. 1e+03;
Matches 15; Conservative 0; Mismatches 2; Indels
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/note= "OTHER= 2' methoxyethyl (2' MOE) nu
Cytidine nucleobases are 5-methylcytidine.
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/label= OTHER= phosphorothioate backbone
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                                                                                                                                                                                            Location/Qualifiers
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Matches
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Gaps

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The present invention relates to antisense compounds targeted to a nucleic acids encoding human and mouse complement component C3. The antisense compound comprises an antisense oligonucleotide that specifically hybridises with the nucleic acid and inhibits the expression
diagnosing a disease associated with LRH1, such as breast cancer, dyslipidaemia, atherosclerosis, low HDL (high density lipoprotein), high LDL (low density lipoprotein), hypercholesterolaemia, gall stones, triglyceridaemia, obesity, hepatitis B virus-mediated acute or chronic hepatitis, as well as hepatocellular carcinoma or a condition associated with aromatase activity. Accordingly, these compositions exhibit cycostatic, antilihaemic, antierrisoslerotic, anorectic, hepatotropic, litholytic, antiinflammatory and virucidal activities. This oligonucleotide sequence is an antisense DNA oligo used to modulate the expression of the human LRH1 protein of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Antisense therapy; mouse; complement component C3; autoimmune disorder; multiple sclerosis; infection; atherosclerosis; neuroprotective; antiarteriosclerotic; antimicrobial; antiinflammatory; cytostatic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /note= "This oligonucleotide has a phosphorothioate backbone and 2'-methyoxyethyl (2'-MOE) wings at the 5' and 3' ends, which are 5 nucleotides in length at each end. All cytidine residues are 5-methylcytidines"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New antisense compound targeted to a nucleic acid molecule encoding complement component C3, useful for treating multiple sclerosis, an
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mouse complement component C3 DNA, antisense oligonucleotide #61.
                                                                                                                                                                                                                                                                                                    Length 20;
                                                                                                                                                                                                                                                                                                                                                Indels
                                                                                                                                                                                                                                                        Sequence 20 BP; 1 A; 6 C; 6 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                             5
                                                                                                                                                                                                                                                                                                    Score 13.8; DB 1;
                                                                                                                                                                                                                                                                                                                           1e+03;
                                                                                                                                                                                                                                                                                                                                                  0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 16; SEQ ID NO 173; 74pp; English.
                                                                                                                                                                                                                                                                                                                             Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /mod_base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                869
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     infection or atherosclerosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        23-OCT-2001; 2001US-00001076
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            18-AUG-2003; 2003US-00642802
                                                                                                                                                                                                                                                                                                                                                                                                                                       m
                                                                                                                                                                                                                                                                                                           0.8%;
                                                                                                                                                                                                                                                                                                                               88.2%;
                                                                                                                                                                                                                                                                                                                                                                                              853 GACAAGGACCTGAAGCA
                                                                                                                                                                                                                                                                                                                                                                                                                                       19 GACAGGGCCCTGAAGCA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADK12315 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ൯
                                                                                                                                                                                                                                                                                        Query Match
Best Local Similarity 88.27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                phosphorothioate; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Graham MJ, Watt AT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2004-225730/21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (GRAH/) GRAHAM M J. (WATT/) WATT A T.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            US2004043956-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (WATT/) WATT A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Key
modified base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           20-MAY-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mus musculus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     04-MAR-2004.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADK12315;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              RESULT 1629
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADK12315
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ð
                                                                                                                                                                                                                                                                                                                                                                                                                                            qq
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     This invention relates to novel antisense compounds useful for modulating the expression of liver related homologue-1 (LRH1) and splice variants thereof. Specifically, it refers to compositions 8-30 nucleobases in length that target a portion of an active site on the nucleocases in molecule encoding LRH1 (also known as NRSA2). LRH1 is a monomeric orphan nuclear receptor protein that functions as a tissue specific transcription factor. The present invention describes antisense oligonucleotides that comprise at least one modified internucleoside linkage, a phosphorothioate linkage, at least one modified supar moiety, a 2'-0-methoxyethyl (2' MOB) and at least one modified supar moiety, methylcytidine. These antisense compounds are useful for treating or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New antisense oligonucleotides targeted to a nucleic acid encoding liver related homologue-1 (LRH1), useful for treating breast cancer, dyslipidemia, atherosclerosis, hypercholesterolemia, or hepatitis.
                                                                                                                                                                                                                                                                                                                                                                           phosphorothioate, 2' MOE, breast cancer, dyslipidaemia, atherosclerosis, low HDL, high density lipoprotein, high LDL, hypercholesterolaemia, gall stone, triglyceridaemia, obesity, hepatitis,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /mod_base= OTHER
/nore= "OTHER= 2' methoxyethyl (2' MOE) nucleotides. All
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /note = "OTHER = 2' methoxyethyl (2' MOE) nucleotides. All
                                                                                                                                                                                                                                                                                                             Antisense DNA oligo used to modulate human LRH1 expression SeqID 147.
                                                                                                                                                                                                                                                                                                                                                                                                                                            hepatocellular carcinoma; arcmatage; gytostatic; antilipaemic; antiarteriosclerotic; anoractic; hepatotropic; litholytic; antiinflammatory; virucidal.
                                                                                                                                                                                                                                                                                                                                                           human; ss; liver related homologue-1; LRH1; NR5A2; antisense;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   cytidine nucleobases are 5-methylcytidine."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 cytidine nucleobases are 5-methylcytidine.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             label= OTHER= phosphorothioate backbone
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Example 15; SEQ ID NO 147; 909pp; English
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                             1448 AACATCCATTCTTCCTC 1464
                                                                                                                                                                                        BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     01-JUL-2002; 2002US-0392813P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           01-JUL-2003; 2003WO-US020865.
                                                                                                                                                                                   ADJ15597 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ...20
/*tag= b
                                                       AACATCCACTCTGCCTC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                *tag= a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            U
                                                                                                                                                                                                                                                                               (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (PHAA ) PHARMACIA CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /*tag=
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modified base

08-JAN-2004

Kane CD;

modified_base

sapiens.

Homo

20-MAY-2004

ADJ15597;

ADJ15597

modified base

and CPRB proteins. The invention is useful for CPRA and CPRB protein

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of complement component C3 in cells. The antisense oligonucleotide is a chimeric oligonucleotide. The antisense oligonucleotide comprises at least one modified internucleoside linkage, preferably a phosphorothioate linkage. It also comprises at least one modified sugar moiety, preferably a 2'-0-methoxyethyl (2'-MOE) sugar moiety. The antisense oligonucleotide further comprises at least one modified nucleobase, preferably a 5-methylcytosine. The antisense oligonucleotides are useful for the treatment of diseases such as autoimmune disorders e.g. multiple sclerosis, infections, and atherosclerosis. The present sequence represents an antisense oligonucleotide used in the examples of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New isolated nucleic acid encoding a CPRA protein, used to increase production of dicarboxylic acid, for use in chemical products including plastics, adhesives, and fragrances.
                                                                                                                                                                                                                      Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Cornett CA;
                                                                                                                                                                                                                                                                                                                                                                                                                                        CYP gene; CPRA protein; CPRB protein; dicarboxylic acid; plastic; adhesive; fragrance; PCR; primer; ss.
                                                                                                                                                                                                                     ..
                                                                                                                                                                                    0.8%; Score 13.8; DB 1; Length 20;
88.2%; Pred. No. 1e+03;
Live 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Madduri KM,
                                                                                                                                                         Sequence 20 BP; 3 A; 9 C; 3 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                           Candida tropicalis CYP gene PCR primer #43.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Eirich LD, Eshoo M,
Loper JC, Gleeson M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example 11; SEQ ID NO 47; 195pp; English.
                                                                                                                                                                                                                                            407
                                                                                                                                                                                                                                                                                                                                      BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 01-MAY-1998; 98US-0083798P.
05-OCT-1998; 98US-0103099P.
10-MAR-1999; 99US-00305620.
12-OCT-2001; 2001US-00976800.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         03-APR-2003; 2003US-00405660.
                                                                                                                                                                                                                                         TCGGATGAGGTGCAGTC
                                                                                                                                                                                                                                                                  rcadardaderecadec
                                                                                                                                                                                                                                                                                                                                  ADJ26672 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Craft DL,
Tang M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ESHOO M.
MADDURI K M.
CORNETT C A.
BRENNER A A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CR.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CRAFT D L.
EIRICH L D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2004-088917/09
                                                                                                                                                                                                  Local Similarity
les 15; Conserv
                                                                                                                                     present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Candida tropicalis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TANG M.
LOPER J C.
GLEESON M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      US2003186411-Al
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WILSON
                                                                                                                                                                                                                                                                                                                                                                                   20-MAY-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Wilson CR,
Brenner AA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                32-OCT-2003
                                                                                                                                                                                                                                         391
                                                                                                                                                                                                                                                                   20
                                                                                                                                                                                                                                                                                                                                                          ADJ26672;
                                                                                                                                                                                      Query Match
                                                                                                                                                                                                                                                                                                       RESULT 1630
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (WILS/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (CORN/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (TANG/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (GLEE/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (CRAF/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Wilson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (EIRI/
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (MADD/
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (ESHO/
                                                                                                                                                                                                             Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                      CYP
                                                                                                                                                                                                                                                                                                                       ADJ26672
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The invention comprises CYP genes from Candida tropicalis which encode

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           production, the DNA and protein sequences are useful for increasing production of dicarboxylic acid in chemical products, such as: plastics, adhesives and fragrances. The present DNA sequence represents a PCR primer that was used in an example of the invention.
                                                                                                                                         Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    treating
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New isolated nucleic acid molecules useful for diagnosing or treati
asthma or bronchial hyperresponsiveness, or other diseases such as
obesity or inflammatory bowel disease.
                                                                                                                                        .;
0
                                                                                                         Score 13.8; DB 1; Length 20;
Pred. No. 1e+03;
0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Del Mastro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0.8%; Score 13.8; DB 1; Length 20; 88.2%; Pred. No. 1e+03;
                                                                                                                                     Indels
                                                                                                                                                                                                                                                                                                                                                                 asthma; bronchial hyperresponsiveness; obesity; inflammatory bowel disease; human; gene 216; ss; primer.
                                                                             Sequence 20 BP; 7 A; 3 C; 9 G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 20 BP; 4 A; 4 C; 7 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Dupuis J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example 10; SEQ ID NO 154; 485pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Eerdewegh PV,
Pandit S;
                                                                                                                                                                1026
                                                                                                                                                                                         18
                                                                                                                                                                                                                                                            BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   13-APR-1999; 99US-0129391P.
13-APR-2000; 2000US-00548797.
13-APR-2001; 2001US-00834597.
                                                                                                           0.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         19-APR-2002; 2002US-00126022
                                                                                                                                                            1010 AGAGGGGAGAGCTCAAG
                                                                                                                                                                                       2 AGAGGGCAGGGCTCAAG
                                                                                                                                                                                                                                                          ADL81342 standard; DNA; 20
                                                                                                                                                                                                                                                                                                              (first entry)
                                                                                                     Query Match 0.8
Best Local Similarity 88.2
Matches 15, Conservative
                                                                                                                                                                                                                                                                                                                                        Gene 216 SSCP primer #92.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DUPUIS J.
DEL MASTRO R G.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    LITTLE R D.
EERDEWEGH P V.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Little RD,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2004-142647/14.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Allen K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SIMON J.
ALLEN K.
PANDIT S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                     US2004023215-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         KEITH T.
                                                                                                                                                                                                                                                                                                                                                                                                          Homo sapiens
                                                                                                                                                                                                                                                                                                              20-MAY-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                               05-FEB-2004.
                                                                                                                                                                                                                                                                                     ADL81342;
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Simon J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (DMAS/)
(SIMO/)
(ALLE/)
(PAND/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (LITT/)
(EERD/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       KEIT/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DUPU/)
                                                                                                                                                                                                                               RESULT 1631
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888888888
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15;

Matches

19

g ð

RESULT 1632

20-MAY-2004

ADL81282

05-FEB-2004

Homo

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This invention relates to a novel primers useful for synthesising full length cDNA molecules that encode human proteins. Specifically, it refers to secretory or membrane proteins that are potential therapeutic agents/ target molecules in the field of medicine, and in particular genes encoding proteins that are associated with signal transduction.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     glycoproteins and transcription. The present invention describes a method for efficiently cloning a full length human cDNA from both the 5' and 3' ends using the oligo-capping method. This oligonucleotide sequence is a human clone specific PCR primer used in an exemplification of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human; NOVX; PCR; ss; congenital heart defect; cardiomyopathy; atherosclerosis; hypertension; pulmonary stenosis; scleroderma; adenocarcinoma; haemophilia; graft-versus-host disease; cancer; neurodegenerative disorder; Alzheimer's disease; Parkinson's disease; multiple sclerosis; diabetes; obesity; bronchial asthma; acquired immunodeficiency syndrome; AIDS; Crohn's disease;
                                                                        Clone specific PCR primer to amplify human full length cDNA SeqID 4416.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New oligonucleotide primers (830 cDNAs) useful for synthesizing full
                                                                                                           human; medicine; signal transduction; glycoprotein; transcription;
                                                                                                                                                                                                                                                                                                                                                                                                                  Hayashi K, Ishii S, Kawai Y;
K, Kojima S, Otsuki T, Koga
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0.8%; Score 13.8; DB 1; Length 20; 38.2%; Pred. No. 1e+03; ve 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 20 BP; 7 A; 3 C; 8 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example 18; SEQ ID NO 4416; 1340pp; English.
                                                                                                                               oligo-capping method; ss; PCR; primer
                                                                                                                                                                                                                                                                                                                                                                                                                     Isogai T,
a T, Nagai
                                                                                                                                                                                                                                                                                                                                                                                    (REAS-) RES ASSOC BIOTECHNOLOGY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        BP.
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11-7AN-2000; 2000JP-00118774.
02-MAY-2000; 2000JP-00183865.
07-JUL-2000; 2000EP-00114089.
                                                                                                                                                                                                                                                                     07-JUL-2000; 2003EP-00025638.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    88.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      19 TGGACAGGAATGCAGAG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TGGACAGGCAAGCAGAG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADM93686 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human NOVX PCR primer #24.
                                                 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Local Similarity 88.2
Les 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                           Wakamatsu A, Sugiyama
                                                                                                                                                                                                                                                                                                                                                                                                                       Nishikawa T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2004-204755/20.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CDNAS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              length human
                                                                                                                                                                                                   EP1396543-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           01-JUL-2004
                                               20-MAY-2004
                                                                                                                                                                                                                                       10-MAR-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADM93686;
                 ADL32383;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          RESULT 1634
                                                                                                                                                                                                                                                                                                                                                                                                                           Ota T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Matches
                                                                                                                                                                       Homo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADM93686
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The invention relates to an isolated nucleic acid molecule, or a set of nucleic acid molecules each given in the specification. The composition and methods are useful in diagnosing or treating asthma or bronchial hyperresponsiveness, and other diseases such as obesity or inflammatory bowel disease. The present sequence is used in the exemplification of the
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     Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New isolated nucleic acid molecules useful for diagnosing or treasthma or bronchial hyperresponsiveness, or other diseases such obesity or inflammatory bowel disease.
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       Indels
                                                                                                                                                                                                                                                                                             asthma; bronchial hyperresponsiveness; obesity; inflammatory bowel disease; human; gene 216; ss; primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 20 BP; 5 A; 7 C; 4 G; 4 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 13.8; DB 1
Pred. No. 1e+03;
); Mismatches
        Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example 10; SEQ ID NO 94; 485pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Eerdewegh PV,
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                                          554
                                                                                                                                                             BP.
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2000US-00548797.
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                                          538 CCCATCTTGACAAGCC
                                                                           CCCTTCTGTGACAAGCC
                                                                                                                                                              ADL81282 standard; DNA; 20
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                                                                                                                                                                                                                                (first entry)
                                                                                                                                                                                                                                                               Gene 216 SSCP primer #32
          Conservative
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LITTLE R D.
EERDEWEGH P V.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DUPUIS J.
DEL MASTRO R G.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Little RD,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2004-142647/14.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   present invention.
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ALLEN K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PANDIT S.
                                                                                                                                                                                                                                                                                                                                                                                  US2004023215-A1.
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13-APR-2000;
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Keith T, Simon J,

(DMAS/) (SIMO/) (ALLE/) (PAND/)

KEIT/) (EERD/) (DUPU/)

Ή,

0;

Gaps

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BP.

ADL32383 standard; DNA; 20

RESULT 1633

ADL32383

15;

Matches

N 538

g à

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infectious disease; anorexia; immune disorder; primer.
                                                                      22-OCT-2001; 2001US-00035568.
05-NOV-2001; 2001US-0338626P.
05-NOV-2001; 2001US-0345398P.
09-NOV-2001; 2001US-0345398P.
15-NOV-2001; 2001US-034513P.
15-NOV-2001; 2001US-0335112P.
28-NOV-2001; 2001US-033312P.
29-NOV-2001; 2001US-0334300P.
04-DEC-2001; 2001US-0334300P.
05-FEB-2002; 2001US-0336576P.
05-FEB-2002; 2002US-0380968P.
                                                                                                                                                                          02-JUL-2002; 2002US-0393148P.
02-JUL-2002; 2002US-0393262P.
                                                         05-NOV-2002; 2002US-002B7971
                                                                                                                                                                                         06-AUG-2002; 2002US-0401479P.
                                                                                                                                                                                                       07-AUG-2002; 2002US-0401593P
07-AUG-2002; 2002US-0401695P
26-AUG-2002; 2002US-0406181P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PENA C E A.
PEYMAN J A.
RASTELLI L.
RIEGER D K.
ROTHENBERG M E.
                                                                                                                                                                                                                                    ALSOBROOK J P.
ALVAREZ E.
ANDERSON D W.
                                                                                                                                                                                                                                                                                                                                                                                                        LI L.
MACDOUGALL J R.
                                                                                                                                                                                                                                                                                                                                                                                  KEKUDA R.
KHRAMTSOV N V.
                                                                                                                                                                                                                                                                                                                                                                                                                     MALYANKAR U M.
                                                                                                                                                                                                                                                                                                                         ETTENBERG S.
GANGOLLI E A.
GERLACH V.
                                                                                                                                                                                                                                                                                                                                                                                                                            MAZUR A.
MCQUEENEY K.
MEZES P S.
MILLER C E.
MILLET I.
MISHRA V.
                                                                                                                                                                                                                                                         BARON M.
BOLDOG F L.
BURGESS C E.
CASMAN S J.
CHAPOVAL A.
DHANABAL M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PADIGARU M.
PATTURAJAN M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SHENOY S G.
SHIMKETS R A.
                                                                                                                                                                                                                                                                                                     EDINGER S R.
                                                                                                                                                                                                                                                                                                           ELLERMAN K.
                                                                                                                                                                                                                                                                                                                                                                                                 LEPLEY D M.
                                                                                                                                                                                                                                                                                                                                               GORMAN L.
GROSSE W M.
                                                                                                                                                                                                                                                                                                                                                             GUO X.
HACKETT C.
                            US2004067882-A1
               Homo sapiens
                                            08-APR-2004.
                                                                                                                                                                                                                                    (ALSO/)
                                                                                                                                                                                                                                           (ALVA/)
                                                                                                                                                                                                                                                                                            (DHAN/)
(EDIN/)
(EISE/)
(ELLE/)
                                                                                                                                                                                                                                                         (BARO/)
(BOLD/)
(BURG/)
                                                                                                                                                                                                                                                                              (CASM/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (ROTH/)
(SHEN/)
(SHIM/)
                                                                                                                                                                                                                                                                                                                                                     (GROS/)
                                                                                                                                                                                                                                                                                                                          ETTE/
                                                                                                                                                                                                                                                                                                                                (GANG)
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                                                                                                                                                                                                                                                                                                                                                                   (HACK/
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(KHRA/
                                                                                                                                                                                                                                                                                                                                                                                                      (LILL/
                                                                                                                                                                                                                                                                                                                                                                                                                     (MALY/
                                                                                                                                                                                                                                                                                                                                                                                                                                   (MCQU/
                                                                                                                                                                                                                                                                                                                                                                                                                                                        (MILL/
(MISH/
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PADI/
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (PATT/)
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RAST/
                                                                                                                                                                                                                                                                                                                                               GORM/
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The invention relates to human NOVX polypeptides and the polynucleotides are useful for encoding them. The NOVX polypeptides and polynucleotides are useful for determining the presence of or predisposition to a disease associated with altered levels of the sequences in a mammalian subject, and for treating or preventing a pathology associated with NOVX. The polypeptides, polynucleotides and antibodies that bind immunospecifically to the polypeptides are useful for treating or preventing disorders or syndromes such as congenital heart defects, cardiomyopathy, atherosclerosis, hypertension, pulmonary stenosis, scleroderma, atherosclerosis, hypertension, pulmonary stenosis, scleroderma, andenocarcinoma, haemophilia, graft-versus-host disease, cancer, neurodegenerative disorders, Alzhaimer's disease, parkinson's disease, immunodeficiency syndrome (ALDS), crohn's disease, preferents a PCR primer used to amplify a human NOVX polynucleotide of the invention. Note: The sequence data for this patent is also available from USPPO at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ·;
                                                                                                                                                                                                                            Alsobrook JP, Alvarez E, Anderson DW, Baron M, Boldog FL; Burgess CE, Casman SJ, Chapoval A, Dhanabal M, Edinger SR, Eisen A; Ellerman K, Ettenberg S, Gangolli EA, Gerlach V, Gorman L; Grosse WM, Guo X, Hackett C, Ji W, Kekuda R, Khramtsov NV. Lepley DM, Li L, Macdougall JR, Malyankar UM, Mazur A, Mcqueney K; Mazes PS, Miller CE, Millet I, Mishra V, Padigaru M, Patturajan M; Pena CEA, Peyman JA, Rastelli L, Rieger DK, Rothenberg ME; Shenoy SG, Shimkets RA, Smithson G, Spaderna SK, Starling G; Spytek KA, Stone DJ, Tchernev VT, Twomlow N, Vernet CAM; Zerhusen BD, Voss EZ, Zhong M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   chimeric; antisense oligonuclectide; phosphorothioate; human; microsomal prostaglandin E2 synthase; mPGES-1; mPGES-1 inhibitor; microsomal prostaglandin E2 synthase inhibitor; cytostatic; antidiabetic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Novel isolated NOVX polypeptide useful treating or preventing disorders or syndromes such as Alzheimer's disease, Parkinson's disease, multiple sclerosis, diabetes, obesity, cancer, bronchial asthma, Crohn's disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human mPGES-1 chimeric antisense oligonucleotide SEQ ID NO:977.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  / Match 0.8%; Score 13.8; DB 1; Length 20; Local Similarity 88.2%; Pred. No. 1e+03; nes 15; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 20 BP; 10 A; 3 C; 5 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example C; SEQ ID NO 318; 330pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          seqdata.uspto.gov/sequence.html.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1240 TICATCITCCGTATCIT 1256
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    BP.
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 SMITHSON G.
SPADERNA S K.
STARLING G.
SPYTEK K.
STONE D J.
                                                                                                                                  VERNET C A M.
ZERHUSEN B D.
VOSS E Z.
                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2004-355303/33.
                                                                                                                 TWOMLOW N.
                                                                                                                                                                                            ZHONG M.
                                                                                                               (TWOM/)
(VERN/)
(ZERH/)
(VOSS/)
(ZHON/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADM14790;
                                                                         (STON/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
                                     (STAR/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RESULT 1635
ADM14790/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Matches
원
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19 GAGGATGCCCTGAGACA 3

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The present sequence represents a chimeric antisense oligonucleotide cargeted to human microsomal prostaglandin E2 synthase (mPGES-1). The human mPGES-1 gene is located on chromosome 9, more specifically to human mpGES-1 gene is located on chromosome 9, more specifically to present invention also describes: (1) antisense compounds, aggas.). Which specifically hybridise with the nucleic acid encoding mPGES-1, which specifically hybridise with the nucleic acid mPGES-1 and inhibits its expression; (2) a method of inhibiting the expression of inhibits its expression; (2) a method of inhibiting the expression of confition associated with mPGES-1. MPGES-1 chimeric antisense oligonucleotides and antisense compounds have cytostatic, antiinflammatory, neuroprotective, nootropic, antianthatic, vasotropic, antiinflammatory, neuroprotective, nootropic, antiarthritic, vasotropic, ophthalmological, immunomodulatory and cardiovascular activities, and can be used for preparing a composition for treating a disease or condition associated with mPGES-1 e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer, ischaemia or reperfusion injury, or ophthalmic, immunological, cardiovascular or neurological disorder.
             neuroprotective; nootropic; antiarthritic; vasotropic; ophthalmological; immunomodulatory; cardiovascular; gene therapy; inflammation; alzheimer; disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder; cardiovascular disorder; neurological disorder;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New antisense compound, having a sequence targeted to a nucleic acid encoding mPGES-1, useful for preparing a composition for treating e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer or
                                                                                                                                                                                                                                                    'note= "phosphorothioate linkages and all cytidine
cardiant; neuroprotective; antiinflammatory;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 20 BP; 2 A; 7 C; 6 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                        residues are 5-methylcytidines
                                                                                                                                                                                                                                                                                                                '*tag= a
'mod_base= OTHER
'note= "2'-O-methocyethyls"
                                                                                                                                                                                                                                                                                                                                                                                                                       /note= "2'-0-methoxyethyls"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 4; SEQ ID NO 977; 132pp; English
                                                                                                                                                                                    location/Qualifiers
                                                                                                                                                                                                                                        /mod_base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                      base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  25-SEP-2003; 2003WO-US030374
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                                                                                                                                                                                                                                                                                                                                                                                       /*tag=
                                                                                                                                                                                                          .20
                                                                                                                                                                                                                         *tag=
                                                                                                                                                                                                                                                                                                                                                                                                        /mod
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         immunomodulator;
                                                                                                                                                                                                                                                                                                                                                                                                                                                               WO2004028458-A2
                                                                                                                                                                                                                                                                                                   modified base
                                                                                                                                                                                                                                                                                                                                                                      modified base
                                                                                                                                                                                                          modified base
                                                                                                                                     sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  08-APR-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gierse JK;
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                                                                                                                                                      Synthetic
                                                                                                                                       Ношо
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The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin E2 synthase (mPGES-1). The human mPGES-1 gene is located on chromosome 9, more specifically to 9q34.3. The present invention also describes: (1) antisense compounds, having a sequence comprising 8-30 bp targeted to a nucleic acid encoding mPCES-1, which specifically hybridise with the nucleic acid mPGES-1 and inhibits its expression; (2) a method of inhibiting the expression of inhibits in cells or tissues; and (3) a method of treating an animal having a disease or condition associated with mPGES-1. MPGES-1 chimeric antisense oligonucleotides and antisense compounds have cytostatic, antidiabetic, immunomodulator, cardiant, neuroprotective,
                                                                                                                                 chimeric; antisense oligonucleotide; phosphorothioate; human; microsomal prostaglandin E2 synthase; mPGES-1; mPGES-1 inhibitor; microsomal prostaglandin E2 synthase inhibitor; cytostatic; antidiabetic; immunomodulator; cardiant; neuroprotective; antiinflammatory; neuroprotective; notiropic; ophthalmological;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New antisense compound, having a sequence targeted to a nucleic acid encoding mPGES-1, useful for preparing a composition for treating e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer or
                                                                                                                                                                                                                                                                                                                                                                                                                   /note= "phosphorothioate linkages and all cytidine residues are 5-methylcytidines"
                                                                                                                                                                                                                     immunomodulatory; cardiovascular; gene therapy; inflammation; Alzheimer's disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder; cardiovascular disorder; neurological disorder; ss.
                                                                                                    Human mPGES-1 chimeric antisense oligonucleotide SEQ ID NO:1120
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /*tag= c
/mod_base= OTHER
/note= "2'-0-methoxyethyls"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /note= "2'-0-methocyethyls"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 4; SEQ ID NO 1120; 132pp; English.
                                                                                                                                                                                                                                                                                                                                                        Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /*tag= a
/mod_base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                        base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          25-SEP-2003; 2003WO-US030374.
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             ADM14933 standard; DNA; 20
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                                                                                (first entry)
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                                                                                                                                                                                                                                                                                                                                                               Key
modified_base
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                                                                                                                                                                                                                                                                                                               sapiens.
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                                                                                01-JUL-2004
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                                                                                                                                                                                                                                                                                                                               Synthetic.
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                                               ADM14933;
                                                                                                                                                                                                                                                                                                               Homo
ADM14933
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Gaps

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Indels

5

0.8%; Score 13.8; DB 1; Length 20; 88.2%; Pred. No. 1e+03;

Pred. No. 1e+03;); Mismatches

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Query Match 0.8 Best Local Similarity 88.2 Matches 15; Conservative

10017621-3sl.rng

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antiinflammatory, neuroprotective, nootropic, antiarthritic, vasotropic, ophthalmological, immunomodulatory and cardiovascular activities, and can be used as mPGES-1 inhibitors and in gene therapy. The antisense compound can be used for preparing a composition for treating a disease or condition associated with mPGES-1 e.g., inflammation, Alzheiner's disease, arthritis, diabetes, cancer, ischaemia or reperfusion injury, or ophthalmic, immunological, cardiovascular or neurological disorder.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Novel single or multiple target oligonuclectide anti-sense to e.g. initiation codon, intron of respiratory disease-relevant gene e.g. CCR1, RANTES, MCP4, useful for prophylaxis or treating respiratory disease e.g.
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human; ss; interleukin-4 receptor; IL-4; interleukin-5 receptor; IL-5; CCR1; Ectaxin-1; RANTES; MCP4; CD23; ICAM, VCAM; tryptase a; tryptase b; PDE4 A; PDE4 C; PDE4 C; PDE4 C; PDE4 D; respiratory disease; lung disease; hyper-responsiveness; adenosine; adenosine A receptor; asthma; lung allergy; inflammation; inflammatory disease, alivay inflammation; allergy; impeded respiration; cystic fibrosis; CF; chronic obstructive pulmonary disease; CODP; allergic rhinitis; acute respiratory distress syndrome; pulmonary hypertension; lung inflammation; bronchitis; airway obstruction; bronchoconstriction.
                                                                                                                                                                                                 Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The invention relates to oligonucleotides anti-sense to an initiation codon, coding region, 5' or 3' intron-exon junction, intron or region
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                                                                                                                                                            0.8%; Score 13.8; DB 1; Length 20; 88.2%; Pred. No. 1e+03; Ive 0; Mismatches 2; Indels
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                                                                                                                         Sequence 20 BP; 2 A; 8 C; 5 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Aguilar D,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 2; SEQ ID NO 2249; 174pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sandrasagra A, Tang L,
n S, Lu H, Cong H;
                                                                                                                                                                                                                          GAGGATGCCATGAAACA 1451
                                                                                                                                                  Query Match
Best Local Similarity 88.2%;
Matches 15; Conservative
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23-APR-2002; 2002WO-US013143.
                                                                                                                                                                                                                                                                                                                                                                                                                                 Human oligonucleotide #2149
                                                                                                                                                                                                                                                                                                                                   ADO46783 standard; DNA; 20
                                                                                                                                                                                                                                                      GAGGATGCCCTGAGACA
                                                                                                                                                                                                                                                                                                                                                                                                  15-JUL-2004 (first entry)
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SANDRASAGRA A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SHAHABUDDIN S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2004-293804/27.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AGUILAR D.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CONG H.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MILLER
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                                                                                                                                                                                                                                                                                                      RESULT 1637
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chosen from a gene encoding interleukin (IL)-4 receptor, interleukin (IL)

- receptor, CCR1, CCR3, Eotaxin-1, RANTES, MCP4, CD23, ICAM, VCAM,

- receptor, CCR1, CCR3, Eotaxin-1, RANTES, MCP4, CD23, ICAM, VCAM,

- receptor, CCR1, CCR3, Eotaxin-1, RANTES, MCP4, CD23, ICAM, VCAM,

- also relates to a method of screening a candidate compound that binds to

- come or more nucleic acid target(8) or expressed product(8), for the

- come or more nucleic acid target(8) or expressed product(8), for the

- come or more nucleic acid target(8) or expressed product(8), for the

- come or mena encoding interleukin-4 receptor, interleukin-5 receptor,

- come or mRNA encoding interleukin-4 receptor, interleukin-5 receptor,

- come or mRNA encoding interleukin-4 receptor, interleukin-5 receptor,

- come or mRNA encoding interleukin-4 receptor, interleukin-5 receptor,

- come or mRNA encoding interleukin-4 receptor, interleukin-5 receptor,

- come or mRNA encoding interleukin-4 receptor, interleukin-5 receptor,

- come or mRNA encoding interleukin-4 receptor, interleukin-5 receptor,

- come or mRNA encoding interleukin-4 receptor, interleukin-5 receptor,

- come or mRNA encoding interleukin-4 receptor, interleukin-5 receptor,

- come or mRNA encoding interleukin-4 receptor, interleukin-5 receptor,

- come or mRNA encoding interleukin-4 receptor,

- come or 
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2-10 nucleotides of the 5'-end or 3'-end of a nucleic acid target
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 20 BP; 6 A; 4 C; 3 G; 7 T; 0 U; 0 Other;
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23-APR-2002; 2002WO-US013143.
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MILLER S.
SHAHABUDDIN S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NYCE J W.
SANDRASAGRA
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(TANG/)
(AGUI/)
(MILL/)
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(LUHH/)
(CONG/)
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The invention relates to oligonucleotides anti-sense to an initiation codon, coding region, 5' or 3' intron-exon junction, intron or region with 2-10 nucleotides of the 5'-end or 3'-end of a nucleic acid target chosen from a gene encoding interleukin (IL)-4 receptor, interleukin (IL)-4 receptor, interleukin (IL)-5 receptor, CCR1, CCR3, Ectaxin-1, RANTES, MCP4, CD23, ICAM, VCAM, CCR1, come to a method of screening a candidate compound that binds to one or more nucleic acid target(s) or expressed product(s), for the cone or more nucleic acid target(s) or expressed product(s), for the cone or more nucleic acid target(s) or expressed product(s), for the cone or more nucleic acid target(s) or expressed product(s), for the cone or more nucleic acid target(s) or expiratory or lung disease. The congonical conding interleukin-4 receptor, interleukin-5 receptor, configuration or many and a respiratory or lung disease are useful for reading or treating a respiratory or lung disease. The useful for preventing or treating a respiratory or lung disease. The cuseful for preventing or treating a respiratory or lung disease. The cuseful for reventing or treating a respiratory or lung disease. The cuseful for preventing or treating a respiratory or lung disease. The cuseful for reventing or treating a respiratory or lung disease. The cuseful for preventing or treating a respiratory or lung disease. The cuseful for preventing or treating a respiratory or lung disease. The cuseful for preventing or treating a denosine and/or levels of adenosine A month of the contract of the 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     receptor(s), and/or asthma and/or lung allergies associated with inflammation or an inflammatory disease. The respiratory or lung disease is chosen from airway inflammation, allergy, asthma, impeded respiration, cystic fibrosis (CF), chronic obstructive pulmonary disease (COPD),
                                                                                                                                               Novel single or multiple target oligonuclectide anti-sense to e.g. initiation codon, intron of respiratory disease-relevant gene e.g. CCR1, RANTES, MCP4, useful for prophylaxis or treating respiratory disease e.g.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    allergic rhinitis, acute respiratory distress syndrome, pulmonary hypertension, lung inflammation, bronchitis, airway obstruction or bronchoconstriction. This sequence represents an oligonucleotide of the
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0.8%; Score 13.8; DB 1; Length 20; 88.2%; Pred. No. 1e+03; 2; Indels ive 0; Mismatches 2; Indels
              Aguilar D, Miller S;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 20 BP; 6 A; 4 C; 4 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                    Claim 2; SEQ ID NO 308; 174pp; English.
                    Sandrasagra A, Tang L,
n S, Lu H, Cong H;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Murine SAC1 DNA PCR primer #422
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                                                    Shahabuddin S, Lu H,
                                                                                                         WPI; 2004-293804/27
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nes 15; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      US2004081964-A1.
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                          Nyce JW,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       RESULT 1639
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The invention relates to SACI polypeptides and the polynucleotides encoding them. The polynucleotides contain a variation associated with sensing carbohydrates, other sweeteners or ethanol. The invention also sensing carbohydrates, other sweeteners or ethanol. The invention also comprising altering SACI activity in the sample and measuring the activity, a method for analysing a polynucleotide in a biological sample, comprising contacting a polynucleotide in a biological sample, with a probe where the probe hybridises to a SACI polynucleotide to form a comprisation complex and detecting the hybridisation complex, a method of identifying susceptibility to obesity or diabetes comprising conparing to mucleotide sequence of the suspected SACI allele with a wild type cucleotide sequence identifies a sequence variation of the SACI and the wild-type sequence identifies a sequence variation of the SACI cucleotide sequence, and a method of treating or preventing obesity, cucleotide sequence, and a method of treating or preventing obesity, cucleotide sequence, and a method of treating or preventing obesity, cucleotide sequence, and a method of treating or preventing obesity, comprises or alcoholism associated with expression of SACI, comprising administering are useful for screening drugs for inhibition or restoration of gene function as antidiabetic, antiobesity or antialcohol consumption therapies and for identifying sweeteners and alcohols. This seconds in a subject and subject a maplify murine SACI DNA of the function.
                                                                                                                                                                                                                                                                               New isolated polynucleotides for sensing carbohydrates, other sweeteners, or ethanol, useful for screening drugs for inhibition or restoration of gene function as antidiabetic, antiobesity or antialcohol consumption
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
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Glutamine-fructose-6-phosphate amidotransferase; diabetes; ischemia;
                                                                                                                                                                                        Tordoff MG;
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                                                                                                                                                                                            Li X, Reed DR,
De Jong PJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 20 BP; 6 A; 3 C; 8 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Chimeric phosphorothicate oligonuclectide #217.
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); Mismatches
                                                                                                                                                                                                                                                                                                                                                                               Example 12; SEQ ID NO 465; 148pp; English.
                                                                                                                                                                                              Li S,
                                                                                                                                                                                                                 Ohman JD, Chatterjee A,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            88.2%;
                                                                                                                                                                                                  Bachmanov AA, Beauchamp GK,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local Similarity 88.2'
Matches 15; Conservative
BEAUCHAMP G K.
                                                                                                         ROSS D A.
OHMAN J D.
CHATTERJEE A.
                                                                       REED D R.
TORDOFF M G.
                                                                                                                                                                DE JONG P J.
                                                                                                                                                                                                                                                    WPI; 2004-340133/31
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                                       LI S.
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                                                                                                                                               (CHAT/)
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   BACH/)
                                                                                                                                OHMA/)
                                                                       (REED/)
                                                                                                           (ROSS/)
                                                                                            TORD/)
                                     /8817)
                                                        LIXX/
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Gaps .. . 0

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The present invention relates to a compound which specifically hybridizes with a nucleic acid molecule encoding GFAT, and inhibits the expression of GFAT. Specifically claimed are antisense oligonucleotides capable of medulating the expression of GFAT, and which comprise any of the 3063 sequences of 20 base pairs, given in the specification. The compound, composition and methods are useful for treating a disease or condition associated with GFAT, such as a disease or condition, e.g. diabetes, a cardiovascular or neurological disorder, ischemia/reperfusion injury. They are also useful in research and diagnostics for modulating the expression of GFAT. The present sequence represents a chimeric phosphorothoate oligonucleotide with 2' molecular chimeric phosphorothoate oligonucleotide with 2' molecular and a deoxy gap, these
                                                                                                                                                                                                                                                                                                                                                                  New compounds, particularly antisense oligonucleotides targeted to a nucleic acid encoding glutamine-fructose-6-phosphate amidotransferase (GFAT), for treating diabetes, a cardiovascular or neurologic disorder, ischemia/reperfusion injury.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GFAT; Antidiabetic; Cardiant;
Glutamine-fructose-6-phosphate amidotransferase; diabetes; ischemia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0.8%; Score 13.8; DB 1; Length 20; 88.2%; Pred. No. 1e+03; ative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 20 BP; 4 A; 5 C; 3 G; 8 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                oligonucleotides inhibit human GFAT expression.
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                                                             /note= "2-methoxyethyl wing
                                                                                                                        not\bar{e}= "2-methoxyethyl wing"
                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 4; SEQ ID NO 217; 175pp; English.
   Location/Qualifiers
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                                               base= other
                                                                                                            base= other
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Best Local Similarity 88.2
Matches 15; Conservative
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                                                                          17. .20
                                                                                                                                                                                                                                                                                                         Crosby SD;
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                                                                                            /*tag=
                                                                                                                                                                                                                                                                       (PHAA ) PHARMACIA CORP.
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                                               /mod
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 Key
modified_base
                                                                         modified base
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modified_base
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The present invention relates to a compound which specifically hybridizes with a nucleic acid molecule encoding GFAT, and inhibits the expression of GFAT. Specifically claimed are antisense oligonucleotides capable of modulating the expression of GFAT and which comprise any of the 3063 sequences of 20 base pairs, given in the specification. The compound, composition and methods are useful for treating a disease or condition associated with GFAT, such as a disease or condition, e.g. diabetes, a cardiovascular or neurological disorder, ischemia/reperfusion injury. They are also useful in research and diagnostics for modulating the expression of GFAT. The present sequence represents a chimeric phosphorothicate oligonucleotide with 2' MOSE wings and a deoxy gap, these
                                                                                                                                                                                                                                                                                                                                 New compounds, particularly antisense oligonucleotides targeted to a
nucleic acid encoding glutamine-fructose-6-phosphate amidotransferase
(GFAT), for treating diabetes, a cardiovascular or neurologic disorder,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               transplant rejection; immune system; rheumatoid arthritis; lupus; inflammatory bowel disease; multiple sclerosis; HIV; AIDS; ss; primer.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0.8%; Score 13.8; DB 1; Length 20; 88.2%; Pred. No. 1e+03; Live 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 20 BP; 4 A; 5 C; 3 G; 8 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               oligonucleotides inhibit human GFAT expression.
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               /mod_base= other
/note= "2-methoxyethyl wing"
17. .20
                                                                          /mod_base= other
/note= "2-methoxyethyl wing"
                                                                                                                                                                                                                                                                                                                                                                                                           Claim 4; SEQ ID NO 158; 175pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    131 GGATGAAGAAGATCAAA 147
                                                                                                                                                                                                              17-OCT-2002; 2002US-0419268P.
                                                                                                                                                                               02-OCT-2003; 2003WO-US03332.
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                                                                                                                                                                                                                                           (PHAA ) PHARMACIA CORP
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                                            modified base
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24-APR-2003; 2003WO-US012946.

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Matches 15; Conservative
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WPI; 2004-400724/37.
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20-DEC-2002;
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                                                                                                                                                                                                                 rejection, e.g. cardiac or kidney transplant rejection, in an individual rejection, e.g. cardiac or kidney transplant rejection, in an individual comprises detecting the expression level of one or more genes. The methods, system and kits are useful in diagnosing or monitoring transplant rejection, e.g. heart, kidney, liver, pancreas, pancreatic rislet, lung, bone marrow or stem cell transplant rejection, and the method is also useful in assessing the immune status of an individual. The methods are also useful in diagnosing and monitoring diseases that involve the immune system, e.g. rheumatoid arthritis, lupus, inflammatory bowel diseases, multiple sclerosis, HIVAIDS or viral, bacterial or fungal infection. The present sequence represents a primer for a 50 mer oligonucleotide marker for diagnosis and monitoring of allograft rejection and other disorders.
                                                                                                                            Diagnosing or monitoring transplant rejection, e.g. heart, kidney, liver, pancreas, pancreatic islet, lung, bone marrow or stem cell transplant rejection, in an individual, comprises detecting the expression level of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    transplant rejection; immune system; rheumatoid arthritis; lupus;
inflammatory bowel disease; multiple sclerosis; HIV; AIDS; ss; primer.
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                                                                                                                                                                                                          The present invention relates to diagnosing or monitoring transplant
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                                                                       Morris M;
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                                                                        Prentice J,
                                                                                                                                                                                                                                                                                                                                                                                Sequence 20 BP; 2 A; 8 C; 5 G; 5 T; 0 U; 0 Other;
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                                                                          Ly N,
                                                                                                                                                                                      Claim 58; SEQ ID NO 1163; 1762pp; English
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (EXPR-) EXPRESSION DIAGNOSTICS INC.
                                                   (EXPR-) EXPRESSION DIAGNOSTICS INC
                                                                          Woodward R,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                   1625 GAGGCCCCAGCAGGCAG 1641
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ADP11872 standard; DNA; 20 BP.
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20-DEC-2002; 2002US-00325899.
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                  24-APR-2002; 2002US-00131831
                              20-DEC-2002; 2002US-00325899
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                                                                           저,
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                                                                                                            WPI; 2004-400724/37.
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Best Local Similarity
Matches 15; Conserv
                                                                            Fry
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Rosenberg S;
                                                                            Wohlgemuth J,
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                                                                                       Rosenberg S;
                                                                                                                                                                     the genes.
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The present invention relates to diagnosing or monitoring transplant crejection, e.g. cardiac or kidney transplant rejection, in an individual comprises detecting the expression level of one or more genes. The comprises detecting the expression level of one or more genes. The comprises detecting the expression level of one or monitoring comethods, system and kits are useful in diagnosing or monitoring cransplant rejection, e.g. heart, kidney, liver, pancreas, pancreatic islet, lung, bone marrow or stem cell transplant rejection, in an exportansplant rejection or mechanical organ replacement rejection, in an individual. The methods are also useful in diagnosing and monitoring can individual. The methods are also useful in diagnosing and monitoring classases that involve the immune system, e.g. rheumatoral arthritis, compliant infection. The present sequence represents a crital, bacterial or fungal infection. The present sequence represents a crital or fungal infection. The present sequence represents a configurate rejection and other disorders.
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Diagnosing or monitoring transplant rejection, e.g. heart, kidney, liver, pancreas, pancreatic islet, lung, bone marrow or stem cell transplant rejection, in an individual, comprises detecting the expression level of
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inflammatory bowel disease; multiple sclerosis; HIV; AIDS; ss; primer.
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Pred. No. 1e+03;
0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 58; SEQ ID NO 756; 1762pp; English.
                                                                                                                                                                                                                            Claim 58; SEQ ID NO 1881; 1762pp; English.
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2002US-00325899
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The present invention relates to diagnosing or monitoring transplant rejection, e.g. cardiac or kidney transplant rejection, in an individual comprises detecting the expression level of one or more genes. The methods, system and kits are useful in diagnosing or monitoring transplant rejection, e.g. heart, kidney, liver, pancreas, pancreatic islet, lung, bone marrow or stem cell transplant rejection, e.g. heart, kidney, liver, pancreas, pancreatic islet, lung, bone marrow or stem cell transplant rejection, in an individual. The method is also useful in assessing the immune status of an individual. The methods are also useful in diagnosing and monitoring diseases that involve the immune system, e.g. rheumatcid arthritis, lupus, inflammatory bowel diseases, multiple sclerosis, HIV/AIDS or viral, bacterial or fungal infection. The present sequence represents a primer for a 50 mer oligonuclectide marker for diagnosis and monitoring of allograft rejection and other disorders.

Sequence 20 BP; 7 A; 3 C; 9 G; 1 T; 0 U; 0 Other;

Gaps 0; DB 1; Length 20; 2; Indels 0.8%; Score 13.8; DB 1 88.2%; Pred. No. 1e+03; Mismatches . Query Match Best Local Similarity 88.2° Matches 15; Conservative

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336 CGAGGACTTGAAGATGG 352 50 cdaddactggaagaagg

à g

ADN48631 standard; DNA; 20 BP

12-AUG-2004 (first entry)

Human Notch3 DNA antisense oligonucleotide #75.

Human; Notch3; ss; antisense oligonucleotide; phosphorothioate linkage; sugar moiety; 5-methylcytosine; 2'-O-methoxyethyl sugar moiety; 5-methylcytosine hyperproliferative disorder; cancer; cytostatic.

Homo sapiens

JS2004102390-A1

27-MAY-2004

21-NOV-2002; 2002US-00301832.

21-NOV-2002; 2002US-00301832

(ISIS-) ISIS PHARM INC.

Dobie KW;

Freier SM,

WPI; 2004-399720/37.

New compounds, particularly oligonucleotides targeted to a nucleic acid encoding Notch3, useful for treating diseases associated with Notch3, e.g. hyperproliferative disorders.

Example 15; SEQ ID NO 86; 74pp; English.

The invention relates to a compound targeted to a nucleic acid molecule encoding the human Notch3 polypeptide. The compound is an antisense oligonucleotide that specifically hybridises with the nucleic acid and inhibits expression of the polypeptide. The antisense oligonucleotide comprises at least one modified internucleoside linkage i.e. a phosphorothioate linkage, at least one modified sugar moiety, preferably a 2.-O-methoxyethy sugar moiety, or at least one modified nucleobase comprising a 5-methylcytosine. The antisense compounds are useful for modulating the expression of the human Notch3 polypeptide and in preparation of a composition for treating hyperproliferative disorders, e.g. cancer. This sequence represents a human Notch3 DNA antisense

.. 0 Human presynaptic cytomatrix protein, PCLO, proximal SNP PCR primer #84. Gaps 0; 1; Length 20; 2; Indels Sequence 20 BP; 4 A; 9 C; 3 G; 4 T; 0 U; 0 Other; 3; DB 1 le+03; melanoma associated polymorphic variation, presynaptic cytomatrix protein, PCLO, SNP; single nucleotide polymorphism, PCR, primer. Query Match 0.8%; Score 13.8; I Best Local Similarity 88.2%; Pred. No. 1e+C Matches 15; Conservative 0; Mismatches Kammerer gene therapy; human; ss; melanoma; oligonucleotide of the invention. 357 TGATGGGGAGAGTGACC 373 Braun A, ВР 17 rearcecercacracic 1 06-NOV-2003; 2003WO-US035879. 06-NOV-2002; 2002US-0424475P. 23-JUL-2003; 2003US-0489703P. ADO56652 standard; DNA; 20 (first entry) Roth RB, Nelson MR, (SEQU-) SEQUENOM INC. WPI; 2004-411721/38 WO2004044164-A2 Homo sapiens. 06-NOV-2002; 12-AUG-2004 27-MAY-2004. ADO56652; RESULT 1646 ADO56652 S X G à d

The invention relates to a method of identifying a subject at risk of melanoma comprising detecting the presence or absence of one or more polymorphic variations associated with melanoma in a nucleic acid sample from a subject. Preventing melanoma in a subject comprises detecting the presence or absence of one or more polymorphic variations associated with melanoma in a nucleic acid sample from a subject; and administering a melanoma preventative to a subject in need thereof based upon the presence or absence of the one or more polymorphic variations in the nucleic acid sample. The preventative reduces ultraviolet (UV) light exposure to the subject. The methods, nucleic acids, proteins, and compositions are useful for treating melanoma. The present sequence represents a human presynaptic cytomatrix protein, PCLO, proximal PCR

Identifying a subject at risk of melanoma, useful for treating melanoma, comprises detecting the presence or absence of one or more polymorphic variations associated with melanoma in a nucleic acid sample from a

Example 6; Page 102; 295pp; English.

Sequence 20 BP; 4 A; 5 C; 4 G; 7 T; 0 U; 0 Other;

Gaps ·, Query Match

0.8%; Score 13.8; DB 1; Length 20;
Best Local Similarity 88.2%; Pred. No. 1e+03;
Matches 15; Conservative 0; Mismatches 2; Indels

160 ATGACACTCCGAGGTGG 176

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The invention describes a new compound (I) comprises 8-80 nucleobases in length targeted to a nucleic acid molecule encoding adipophilin, where the compound specifically hybridises with the nucleic acid and inhibits the expression of adipophilin. Also described are: a method of inhibiting the expression of adipophilin. Also described are: a method of inhibiting the expression of adipophilin is of a dispophilin is comprising contacting a sesociated with breast cancer-1, comprising administering (I) to the animal so that expression of adipophilin is inhibited; a method of screening for a modulator of adipophilin, comprising contacting a correct animal so that expression of adipophilin, comprising contacting a concern more candidate modulators, and identifying for one or more candidate modulators, and identifying of cone or more candidate modulators, and identifying a disease state, adipophilin; a diagnostic method for identifying a disease state, adipophilin; a diagnostic method for identifying a disease state, comprising identifying the presence of adipophilin in a sample using at least one of the primers selected from a fully defined sequence of 20, 18 cor assay comprising (I). The methods and compositions of the present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New antisense oligonucleotide compounds, useful for diagnosing, preventing and/or treating diseases or conditions associated with aberrant expression or activity of adipophilin, such as atherosclerosis.
                                                                                                                                                                                                                                                                                                                                                                                                                      'note = "OTHER Phosphorothioate backbone. All cytidines
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               'note= "OTHER= 2'-0-Methoxyethyl (2'-MOE) nucleotides"
                                                                                                                                                                                                                                           antiarteriosclerotic; adipophilin modulator; adipophilin; adipophilin inhibitor; breast cancer-1; atherosclerosis; human; adipophilin; antisense oligonucleotide; antisense technology; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (2'-MOE)
                                                                                                                                                                                                             Human adipophilin antisense oligonucleotide segid 43.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /mod_base= OTHER
/note= "OTHER= 2'-O-Methoxyethyl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example 15; SEQ ID NO 43; 36pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                          are 5-methylcytidines"
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/mod_base= OTHER
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19 ATAACACTCCAAGGTGG
                                                                                                       ADO85087 standard; DNA; 20
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15. .20
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modified_base
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                                                                                                                                            ADO85087;
                                                                                         AD085087,
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nucleotides"

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The invention relates to novel antisense compounds targeted to a nucleic acid molecule encoding human Talin to and inhibit its expression. The invention is useful for treating a disease or condition associated with
invention are useful for the diagnosis, prevention and/or treatment of diseases or conditions associated with aberrant expression or activity adipophilin, such as atherosclerosis. This sequence represents a human adipophilin antisense oligonucleotide.
                                                                                                                           Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New compounds, particularly antisense oligonucleotides targeted to
nucleic acid encoding talin, useful for treating muscular, cardiac,
hematologic, or hyperproliferative disorders.
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                                                                                                                                                                                                                                                                                                                                                       Antisense; Talin; muscular disorder; haematologic disorder; cardiac disorder; hyperproliferative disorder; cancer; human;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /*tag= a
/mod_base= OTHER
/note= "2'-methoxyethyl (2'-MOE) nucleotides"
16. .20
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/note= "2'-methoxyethyl (2'-MOE) nucleotides"
                                                                                              Score 13.8; DB 1; Length 20;
Pred. No. 1e+03;
0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                               Human Talin antisense oligonucleotide, ISIS #109176.
                                                                   Sequence 20 BP; 1 A; 7 C; 4 G; 8 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 3; SEQ ID NO 77; 48pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Location/Qualifiers
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                                                                                                                                                     187 GACAAGACCAATGGTGC 203
                                                                                                                                                                                                                                                   BP.
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                                                                                                 0.8%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             30-OCT-2000; 2000US-00702251
                                                                                                                                                                                                                                                   ADP85732 standard; DNA; 20
                                                                                                                                                                                20 GACAAGACCAAAGGGGC
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                                                                                                              Best Local Similarity 88.2
Matches 15; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2004-440384/41.
                                                                                                                                                                                                                                                                                                                                                                                     phosphorothioate;
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                                                                                                                                                                                                                                                                                                       26-AUG-2004
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          10-JUN-2004
                                                                                                                                                                                                                                                                                                                                                                                                                               Synthetic.
                                                                                                                                                                                                                                                                              ADP85732;
                                                                                                 Query Match
                                                                                                                                                                                                                           RESULT 1648
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ID ADP8
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Talin such as a disease or condition e.g. muscular, haematologic, cardiac or hyperproliferative disorder such as cancer. The present sequence is an antisense oilgonucleotide targeted to human Talin DNA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New antisense oligonucleotides for modulating NF-kappaB repressing factor expression, useful for diagnosing, preventing or treating diseases or conditions involving an immune response.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The invention relates to a compound 8-80 nucleobases in length targeted to a nucleic acid molecule encoding NF-kappaB repressing factor (NRF). NF-kappaB (nuclear factor kappaB) is involved in such cellular processes as the immune response, inflammatory response, oncogenesis, apoptosis, cell cycle, differentiation and cell migration. The compound (an antisense oligonucleotide) specifically hybridises with the nucleic acid molecule
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1..20
/*tag= b
/mod_base= OTHER
/note= "Phosphorothioate backbone and all cytidines are
-methylcytidines"
                                                                                                                     Gaps
                                                                                                                                                                                                                                                                                                                                                         Human; ss; antisense; NRF; NF-kappaB repressing factor;
nuclear factor kappaB; immune response; inflammatory response;
oncogenesis; apoptosis; cell cycle; differentiation; cell migration;
                                                                                                                     0;
                                                                                         Length 20;
                                                                                    0.8%; Score 13.8; DB 1; Length 2
88.2%; Pred. No. 1e+03;
ive 0; Mismatches 2; Indels
                                                        Sequence 20 BP; 5 A; 7 C; 4 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /note= "2'-methoxyethyl residue" 16..20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /note= "2'-methoxyethyl residue"
                                                                                                                                                                                                                                                                                                                                Human NRF antisense oligonucleotide ISIS264076.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 15; SEQ ID NO 82; 61pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Location/Qualifiers
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/mod_base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    'mod base= OTHER
                                                                                                                                          1571 ACTCAGGCAGGCCAGCT 1587
                                                                                                                                                                        20
                                                                                                                                                                                                                                               BP
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                                                                                                                                                                      4 ACTCTGGCAGGCCATCT
                                                                                                                                                                                                                                            ADP74448 standard; DNA; 20
                                                                                                                                                                                                                                                                                                     (first entry)
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                                                                                                Local Similarity 88.2
les 15; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                      chromosome Xq24-25
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                                                                                                                                                                                                                                                                         ADP74448;
                                                                                    Query Match
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encoding NRF (which appears as ADP74371 and comprises nucleotides 469701-489000 of the X chromosome containing the NRF gene at Xq24-25) and inhibits the expression of NRF. Also included are inhibiting the expression of NRF in cells or tissues, screening for a modulator of NRF, a diagnostic method for identifying a disease state, a kit or assay device comprising the above compound, and treating an animal having a disease or condition associated with NRF. The antisense oligonucleotide is useful for inhibiting the expression of NRF in cells or tissues to prevent or treat diseases associated with aberrant NRF expression, such as diseases or conditions involving an immune response. In addition, the compound is used for diagnostics, prophylaxis, or as research reagents or kits. The present sequence represents an antisense oligonucleotide
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /mod_base= OTHER
/note= "OTHER= Phosphorothioate backbone. All cytidines
are 5-methylcytidines"
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/note= "OTHER= 2'-0-Methoxyethyl (2'-MOE) nucleotides"
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                                                                                                                                                                                                                                                                                         Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mouse; Angiopoietin-2; ss; antisense oligonucleotide;
phosphorothicate linkage; 2'-O-methoxyethyl sugar moiety;
5-methylcytosine; hyperproliferative disorder; cancer; cytostatic.
                                                                                                                                                                                                                                                                                         ;
                                                                                                                                                                                                                                                      Score 13.8; DB 1; Length 20;
Pred. No. 1e+03;
0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /mod_base= OTHER
/note= "OTHER= 2'-O-Methoxyethyl (2'-MOE)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Murine Angiopoietin-2 DNA antisense oligonucleotide #6.
                                                                                                                                                                                                                     Sequence 20 BP; 6 A; 5 C; 5 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Location/Qualifiers
                                                                                                                                                                                                                                                                                                                     1182
                                                                                                                                                                                                                                                                                                                                                                                                                             ADQ09470 standard; DNA; 20 BP.
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l Similarity 88.2%;
15; Conservative (
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                                                                                                                                                                                                                                                                                                                  1166 TGGGCTGCATCTTCTAT
                                                                                                                                                                                                                                                                                                                                   20 TGGGCTGCAGCTTCCAT
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                                                                                                                                                                                                                                                                   Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  US2004115640-A1.
                                                                                                                                                                                        targeting NRF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Key
modified_base
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                                                                                                                                                                                                                                                      Query Match
                                                                                                                                                                                                                                                                                                                                                                                               1650
                                                                                                                                                                                                                                                                                    Matches
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Gaps

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Indels Length

DB 1; le+03;

0.8%; Score 13.8; E 18.2%; Pred. No. 1e+0 .ve 0; Mismatches

88.28;

615

BP.

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ADP68684;
                                          Mckay R,
                                             useful
                     1651
                Matches
                      ADP68684/
                       g
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treating an animal having a disease or condition associated with PPAR-alpha. The oligonuclectide compound is useful for preparing a composition for treating hyperproliferative disorder e.g. cancer. This sequence represents a mouse peroxisome proliferator-activated receptor-alpha (PPAR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   cytostatic; gene therapy; PPAR-alpha; peroxisome proliferator-activated receptor-alpha; PPAR-alpha modulator; PPAR-alpha associated disorder; hyperproliferative disorder; mouse; antisense oligonucleotide; antisense technology; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mouse PPAR-alpha antisense oligonucleotide seqid 236.
                                                                                                                                 Sequence 20 BP; 3 A; 5 C; 5 G; 7 T; 0 U; 0 Other;
                                                                                             -alpha) antisense oligonucleotide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             11-DEC-2002; 2002US-00317500.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                11-DEC-2002; 2002US-00317500
                                                                                                                                                                                                                                                                       599 TTGGGAAACTGGAGACC
                                                                                                                                                                                                                                                                                                             17 TTCGGAAACTGCAGACC
                                                                                                                                                                                                                                                                                                                                                                                                                       ADP68800 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
                                                                                                                                                                                                                               15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2004-449378/42.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mckay R, Dobie KW;
                                                                                                                                                                                       Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  US2004115637-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          17-JUN-2004.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            09-SEP-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADP68800;
                                                                                                                                                                                                                                                                                                                                                                                   1652
                                                                                                                                                                                                                                 Matches
                                                                                                                                                                                                                                                                                                                                                                                                         ADP68800
                                                                                                                                                                                                                                                                                                                                                                                     RESULT
                                                                                                                                                                                                                                                                                                                                                                                                                                                 888888888
                                                                                                                                                                                                                                                                            8
                                                                                                                                                                                                                                                                                                                  셤
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The invention describes a compound, having a sequence comprising 8-80 bp targeted to a nucleic acid encoding PPAR-alpha (peroxisome proliferator-activated receptor-alpha), that specifically hybridises with the nucleic acid encoding PPAR-alpha comprising 8601-bp sequence and inhibits expression of PPAR-alpha. Also described are: a method of inhibiting the modulator of PPAR-alpha in cells or tissues; a method of screening for a modulator of PPAR-alpha; a diagnostic method for identifying a disease state; a kit or assay device comprising the compound; and a method of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0
                                                                         The invention relates to a compound targeted to a nucleic acid molecule encoding the human Angiopoietin-2 polypeptide. The compound is an antisense oligonuclectide that specifically hybridises with the nucleic acid and inhibits expression of the polypeptide. The antisense oligonuclectide comprises at least one modified internucleoside linkage i. e. a phosphorothicate linkage at least one modified sugar moiety, preferably a 2'-O-methoxyethyl sugar moiety, or at least one modified nucleobase comprising a 5-methylcytosine. The antisense compounds are useful for modulating the expression of the human Angiopoietin-2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                cytostatic; gene therapy; PPAR-alpha; pportoralpha; PPAR-alpha modulator; peroxisome proliferator-activated receptor-alpha spociated disorder; hyperproliferative disorder; mouse; antisense oligonucleotide; antisense technology; ss.
                                                                                                                                                                                                                                                                       polypeptide and in preparation of a composition for treating hyperproliferative disorders, e.g. cancer. This sequence represents an antisense oligonucleotide targeted to DNA encoding the murine Angiopoietin-2 polypeptide of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           oligonuclectide compound that inhibits expression of PPAR-alpha,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ö
                                                                                                                                                                                                                                                                                                                                                                                                                                0.8%; Score 13.8; DB 1; Length 20; 38.2%; Pred. No. 1e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mouse PPAR-alpha antisense oligonucleotide segid 120.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2;
                                                                                                                                                                                                                                                                                                                                                                                          Sequence 20 BP; 3 A; 7 C; 5 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 16; SEQ ID NO 120; 121pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mismatches
                                            Example 16; SEQ ID NO 106; 102pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ..
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         817 ACGGAGAAGTCCCTCAC 833
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               11-DEC-2002; 2002US-00317500
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    11-DEC-2002; 2002US-00317500
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               19 Acedadedagecrercae 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                        88.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADP68684 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 disorders, e.g. cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 2004-449378/42.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Dobie KW;
                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    US2004115637-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mus musculus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              17-JUN-2004
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
New oligonucleotide compound that inhibits expression of PPAR-alpha, useful for preparing a composition for treating hyperproliferative disorders, e.g. cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     .
0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0.8%; Score 13.8; DB 1; Length 20;
llarity 88.2%; Pred. No. 1e+03;
Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 20 BP; 7 A; 5 C; 5 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                       Example 16; SEQ ID NO 236; 121pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 -alpha) antisense oligonucleotide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Best Local Similarity Matches 15; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
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/mod_base= OTHER
/note= "Phosphorothioate backbone and all cytidines are 5
-methylcytidines"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The invention relates to an oligomeric compound (an antisense oligonucleotide) 8-50 nucleobases in length comprising a sequence complementary to a nucleic acid molecule encoding dual specific phospharatase ( DUST6, phosphorylating Map kinase and extracellular signal related kinase, ERK) appearing as ADP96435. Also included are a composition comprising the oligonucleotide (and a pharmaceutical carrier or diluent) and a method of inhibiting the expression of dual specific phosphatase 6 in cells or tissues comprising contacting the cells or tissues with the antisense oligonucleotide. The oligomeric compound inhibits the expression of dual specific phosphatase 6 by at least 60% and hybridises to nucleobases 369-389, 480-500, 657-677, 713-818, 923-1028, 1196-1216, 12771693, or 1757-1860 in the codding region of $SQ ID NO: 4. The oligomeric compound hybridises to nucleobases 53-195 in the 5'
                                                                                                                                                                                                                                       Human; antisense; ss; dual specific phosphatase 6; DUST6; MAP kinase;
extracellular signal related kinase; ERK; hyperproliferative disorder;
developmental disorder; neural disorder; apoptotic disorder;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New antisense oligonucleotides which inhibit the expression of dual specific phosphatase 6, useful for e.g. treating disease or condition associated with the expression of dual specific phosphatase.
                                                                                                                                                                                                         Human DUSP6 antisense oligonucleotide ISIS103229.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /mod_base= OTHER
/note= "2'-methoxyethyl residue"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /note= "2'-methoxyethyl residue"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Example 15; SEQ ID NO 29; 54pp; English.
                                                                                                                                                                                                                                                                                                                                                        Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /*tag= c
/mod base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Dobie KW;
   615
                             4 Trcggaaacrgcagacc 20
                                                                                                               BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            09-FEB-2004; 2004US-00774888
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            18-JUL-2002; 2002US-00199221
599 TTGGGAAACTGGAGACC
                                                                                                            ADP96460 standard; DNA; 20
                                                                                                                                                                         23-SEP-2004 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                      *tag= b
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  *tag= a
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                                                                                                                                                                                                                                                                                                                                                                       .20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (MONI/) MONIA B P.
(COWS/) COWSERT L M.
(DOBI/) DOBIE K W.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2004-499137/47.
                                                                                                                                                                                                                                                                                          chromosome 12q22-23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               US2004127451-A1
                                                                                                                                                                                                                                                                                                                                                                   modified base
                                                                                                                                                                                                                                                                                                                                                                                                                                                 modified base
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                                                                                                                                                                                                                                                                                                                        Homo sapiens
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                                                                                                                                           ADP96460;
                                                                              RESULT 1653
                                                                                             ADP9646(
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UTR of ADP96435 or to nucleobases 1757-1860 in the 3' UTR of ADP96435. The antisense compound is useful for inhibiting the expression of dual specific phosphatase 6 and for treating a disease or condition associated with the expression of dual specific phosphatase 6 (e.g. a hyperproliferative disorder, developmental disorder, neural disorder or a apoptotic disorder. These may also be used as research reagents and diagnostics, to distinguish between functions of various members of a biological pathway, and in the treatment of a disease or disorder, which can be treated by modulating the expression of dual specific phosphatase 6. The DUST6 gene is located on chromosome 12q22-23. The present sequence is an antisense oligonucleotide targeting DUST6.
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2003 to correct PR field.)
                                                                                                                                                                                                                                                                Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
                                                                                                                                                                                                                                                                0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Detecting and typing human papilloma-virus - using consensus pr
polymerase chain reaction to amplify particular genomic regions
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 .
0
                                                                                                                                                                                                                             0.8%; Score 13.8; DB 1; Length 20;
88.2%; Pred. No. 1e+03;
iive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               HPV6 typing probe (MY12) for use with L1 consensus primers.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 21;
                                                                                                                                                                                                                                                              2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          / Match 0.8%; Score 13.8; DB 1; Length 2 Local Similarity 88.2%; Pred. No. 1.1e+03; les 15; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Papilloma-virus; consensus primer; PCR; probe; ss.
                                                                                                                                                                                            Sequence 20 BP; 7 A; 4 C; 6 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 21 BP; 6 A; 8 C; 1 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Disclosure; Table 5; 33pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1677 CCCCAACTACATCTTCC 1693
                                                                                                                                                                                                                                                                                          445 AAGATCTCCACTGAGGA 461
                                                                                                                                                                                                                                                                                                                         17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Ting Y;
                                                                                                                                                                                                                                                                                                                                                                                                     AAQ03910 standard; DNA; 21 BP.
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89US-00322550.
                                                                                                                                                                                                                                                                                                                       1 AAGATCTCCACTGGGAA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (revised)
(first entry)
                                                                                                                                                                                                                                                           15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Manos MM, Wright DK,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 1990-116005/15.
                                                                                                                                                                                                                                            Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (CETU ) CETUS CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           09-SEP-1988;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         09-SEP-1988;
10-MAR-1989;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 25-MAR-2003
24-AUG-1990
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                    AAQ03910;
                                                                                                                                                                                                                               Query Match
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                                                                                                                                                                                                                                                                                                                                                                     RESULT 1654
                                                                                                                                                                                                                                                           Matches
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RESULT 1655

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Synthetic.

AAQ27035

AAQ27035

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Protein contg. non-A non-B hepatitis antigen fragment - prepd. by culturing transformants transformed by vector contg. base sequence coding specified aminoacid sequences, used for detecting hepatitis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The sequence is that of HPV6 typing probe MY12 for use with L1 consensus primers as part of a simple and rapid assay method for detecting and typing HPV in biological samples. (Updated on 25-MAR-2003 to correct PF field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Detection of genital human papilloma virus - by PCR amplification using defined consensus primer pairs.
                                                                                                                                                                               for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human papilloma virus; amplification; polymerase chain reaction; PCR;
                                                                                                                                                                                                                                                                                                  Gaps
                                                                                                                                                               The present sequence is a primer for a DNA encoding a non-A non-B hepatitis virus (NANBH) or hepatitis C virus (HCV) antigen, useful
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        .
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                                                                                                                                                                                                                                                              Query Match 0.8%; Score 13.8; DB 1; Length 21; Best Local Similarity 88.2%; Pred. No. 1.1e+03; Matches 15; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Wright DK;
                                                                                                                                                                                                                                 Sequence 21 BP; 1 A; 5 C; 13 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 21 BP; 6 A; 8 C; 1 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Manos MM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           consensus primer HPV6 typing probe MY12.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Ting Y,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Disclosure, Page 8; 13pp; English.
                                                                                                                                   Example 1; Page 8; 53pp; Japanese
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                88US-00243486.
89US-00322550.
89WO-US003747.
                                                                                                                                                                                                                                                                                                                                        228 GAGTGGTGGTGGCG 244
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (UYRP ) UNIV ROCHESTER. (HOFF ) HOFFMANN LA ROCHE INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                           BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (revised)
(first entry)
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(GREC ) GREEN CROSS CORP.
                                                                                                                                                                                                      diagnosis or detection
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   detection; assay; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 1994-048082/06.
                                 WPI; 1993-260858/33
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    15-FEB-1991;
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10-MAR-1989;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     29-AUG-1989;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  01-FEB-1994.
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29-JUL-1994
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Synthetic,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAQ56381;
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Matches
                                                                                                                                                                                                                                                                                                                                                                                                                              RESULT 1657
                                                                                                                                                                                                                                                                                                                                                                                                                                                AAQ56381
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     ð
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0;
                                                                                                                                                                                                                                                                                                                                                                                                                                            Hepatitis C virus antigen expressed as recombinant in E.coli - useful for diagnosis of hepatitis C virus infection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The sequences given in AAQ27030-77 are primers. These were used to amplify the claimed hepatitis C virus genes of the invention which could then be inserted into an E. Coli vector. The polypeptides encoded by the vectors were useful as diagnostic reagents for type C hepatitis and they may be produced efficiently by recombinant methods
                                                                                                                                                       E. coli; diagnostic; reagent; type C hepatitis; PCR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            non-A non-B hepatitis virus; NANBH; hepatitis C virus; HCV; antigen; diagnosis; detection; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        .,
0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0.8%; Score 13.8; DB 1; Length 21;
llarity 88.2%; Pred. No. 1.1e+03;
Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 21 BP; 1 A; 5 C; 13 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Primer for hepatitis C virus antigen DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Disclosure, Page 64; 66pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           228 GAGIGGIGGIGGIGGCG 244
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       21
                                                                                                                                                                        polymerase chain reaction; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     91JP-00354708
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                          ВР
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                          AAQ27035 standard; DNA; 21
                                                                                              (first entry)
                                                                                                                                                                                                                                                                                                                                                                                  (TOKU ) TOKUYAMA SODA KK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (SHIM/) SHIMOTONO K.
                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 1992-263663/32.
                                                                                                                                                               vector;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Best Local Similarity
Matches 15, Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Synthetic.
Hepatitis virus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    JP05176774-A.
                                                                                                                                HCV primer P6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      18-DEC-1991;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    18-DEC-1990;
                                                                                                                                                                                                                                               JP04179482-A.
                                                                                                                                                                                                                                                                                                                                                 11-NOV-11990;
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                                                                                                                                                                                                                                                                                                                 11-NOV-1990;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     20-JUL-1993
                                                                                                                                                          Recombinant
                                                                                              21-JAN-1993
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AAV05593;

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8 g RESULT 1656

Query Match

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